CORRECTED VERSION

(19) World Intellectual Property Organization International Bureau





(43) International Publication Date 15 March 2001 (15.03.2001)

PCT

(10) International Publication Number WO 01/018045 A1

(51) International Patent Classification7: C07K 14/00, G01N 33/573

(21) International Application Number: PCT/US00/24700

(22) International Filing Date:

8 September 2000 (08.09.2000)

(25) Filing Language:

English

(26) Publication Language:

English

60/152,753

8 September 1999 (08.09.1999)

(30) Priority Data:

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- (81) Designated States (national): CA, JP, US.
- (84) Designated States (regional): European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE).

Published:

with international search report

(48) Date of publication of this corrected version:

7 November 2002

(15) Information about Correction:

see PCT Gazette No. 45/2002 of 7 November 2002, Section II

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

(54) Title: CRYSTAL STRUCTURE OF A DEACETYLASE AND INHIBITORS THEREOF

(57) Abstract: The present invention provides three-dimensional structural information from the hyperthermophilic bacterium Aquifex aeolicus which is a histone deacetylase-like protein (HDLP). HDLP shares 35.2% amino acid sequence identity with human histone deacetylase (HDAC1). The present invention further provides three-dimensional structural information of HDLP bound by inhibitor molecules. The three-dimensional structural information of the present invention is useful to design, isolate and screen deacetylase inhibitor compounds capable of inhibiting HDLP, HDAC family members and HDLP-related molecules. The invention also relates to nucleic acids encoding a mutant HDLP which facilitates the determination of the three-dimensional structure of HDLP in the presence of a zinc atom.



Crystal Structure of a Deacetylase and Inhibitors Thereof

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This application claims priority of U.S. Provisional Application No. 60/152,753, filed September 8, 1999, the contents of which are hereby incorporated by reference.

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This invention has been made with government support under National Institutes of Health Grant No. RO1 CA-65698. Accordingly, the U.S. Government may have certain rights in the invention.

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Throughout this application, various publications are referenced by author, date and citation. The disclosures of these publications in their entireties are hereby incorporated by reference into this application in order to more fully describe the state of the art as known to those skilled therein as of the date of the invention described and claimed herein.

Introduction

The present invention relates to a histone deacetylase homologue from the hyperthermophilic bacterium Aquifex aeolicus, HDLP (histone deacetylase like protein; also known as AcuCl), which shares 35.2 % sequence identity with human histone deacetylase (HDACl), that can be co-crystallized with an inhibitory ligand, and more particularly, to the detailed crystallographic data obtained from said co-crystallization which is disclosed herein. The invention also relates to methods of using the crystal structure and x-ray crystallographic coordinates of the apo-HDLP and

inhibitor-bound HDLP to design, isolate and screen compounds which bind to and inhibit the active site of HDLP and HDLP-related proteins, such as those proteins belonging to the HDAC family, including HDAC1.

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Background of the Invention

The reversible modification of histones by acetylation is associated with changes in nucleosome conformation and chromatin structure, and plays an important role in the regulation of gene expression (reviewed in Davie and Chadee, 1998, J. Cell Biochem. Suppl. 30-31:203-213). The histone acetylase and deacetylase enzymes that carry out these modifications are involved in many cellular processes such as cell cycle progression and differentiation, and their deregulation is associated with several types of human cancer (reviewed in Kouzarides, 1999, Curr. Opin. Genet. Dev. 9:40-48; Hassig et al., 1997, Chem. Biol. 4:783-789; Fenrick and Heibert, 1998, J. Cell. Biochem. Suppl. 30-31:194-202).

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Recently, several experimental antitumor compounds, such as trichostatin A (TSA), trapoxin, suberoylanilide hydroxamic acid (SAHA), and phenylbutyrate have been shown to act, at least in part, by inhibiting histone deacetylases. Richon et al., 1998, Proc. Natl. Acad. Sci., USA 95:3003-3007; Yoshida et al., 1990, J. Biol. Chem. 265:17174-17179; Kijima et al., 1993, J. Biol. Chem. 268:22429-22435. Additionally, diallyl sulfide and related molecules (Lea et al., 1999, Int. J. Oncol. 2:347-352), oxamflatin (Kim et al., 1999, Oncogene 15:2461-2470), MS-27-275, a synthetic benzamide derivative (Saito et al., 1999, Proc. Natl. Acad. Sci. 96:4592-4597),

butarate derivatives (Lea and Tulsyan, 1995, Anticancer Res. 15:879-883), FR901228 (Nokajima et al., 1998, Exp. Cell Res. 241:126-133), depudecin (Kwon et al., 1998, Proc. Natl. Acad. Sci. USA 95:3356-3361) and m-carboxysinnamic acid bishydroxamide (CBHA; Richon et al., Proc. Natl. Acad. Sci. USA 95:3003-3007) have been shown to inhibit histone deacetylases. In vitro, these compounds can inhibit the growth of fibroblast cells by causing cell cycle arrest in the G1 and G2 phases (Richon et al., 1996, Proc. Natl. Acad. USA 93:5705-5708; Kim et al., 1999, Oncogene 18:2461-2470; Yoshida et al., 1995, Bioessays 17:423-430; Yoshida & Beppu, 1988, Exp. Cell. Res. 177:122-131), and can terminal differentiation and loss to the transforming potential of a variety of transformed cell lines. Richon et al., 1996, Proc. Natl. Acad. Sci. USA 93:5705-5708; Kim et al., 1999, Oncogene 18:2461-2470; Yoshida et al., 1987, Cancer Res. 47:3688-3691. phenylbutyrate is effective in the treatment of acute promyelocytic leukemia in conjunction with retinoic acid. Warrell et al., 1998, J. Natl. Cancer Inst. 90:1621-1625. SAHA is effective in preventing the formation of mammary tumors in rats, and lung tumors in mice. Desai et al., 1999, Proc. AACR 40: abstract #2396; Cohen et al., Cancer Res., submitted.

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Histone deacetylases catalyze the removal of acetyl groups from the e-amino groups of lysine residues clustered near the N-terminus of nucleosomal histones, and this process is associated with transcriptional repression (reviewed in Struhl, 1998, Genes Dev. 12:599-606). Deletion of the yeast histone deacetylase gene, rpd3, or its pharmacological

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inactivation with trichostatin A reduces the transcriptional repression in a subset of promoters, such as those of Ume6-regulated genes. Kadosh & Struhl, 1998, Mol. Cell. Biol. 18:5121-5127. This is accompanied by the increased acetylation of H4 histones in the repressed promoter and its vicinity, but has no effect on histones at promoter distal regions. Kadosh & Struhl, 1998, Mol. Cell. Biol. 18:5121-5127; Rundlett et al., 1998, Nature 392:831-835.

Histone deacetylases are recruited to specific promoters by associating with DNA-binding transcriptional repressors, either directly or through co-repressors that bridge the deacetylase to the transcriptional repressors. For example, the Mad and Ume6 repressors bind to the co-repressor Sin3A (Laherty et al., 1997, Cell 89:349-356; Hassig et al., 1997, Cell 89:341-347; Kadosh & Struhl, 1997, Cell 89:365-371), and the nuclear receptors bind N-CoR and the related SMRT co-repressors. Nagy et al., 1997, Cell 89:373-380; Alland et al, 1997, Nature 387:49-55; Heinzel et al, 1997, Nature 387:43-48.

The deregulation of histone deacetylase recruitment appears to be one of the mechanisms through which these enzymes contribute to tumorigenesis. In acute promyelocytic leukemia (APL), chromosomal translocations fuse the retinoic acid receptor-α (RARα) to either PLZF or to PML. These fusion oncoproteins have aberrant transcriptional repression activity resulting, in part, through the recruitment of a co-repressor and, in turn, HDACs. Grignani et al, 1998, Nature 391:815-818; Lin et al., 1998, Nature 391:811-814. Treatment of PLZF-RARα APL cells with TSA enhances their

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responsiveness to retinoic acid-induced differentiation. Grignani et al, 1998, Nature 391:815-818; Lin et al., 1998, Nature 391:811-814.

5 The histone deacetylases comprise a large family of proteins, conserved from yeast to man, and are divided into two related classes. Class I is characterized by human HDAC1, 2, 3 (Taunton et al., 1996, Science 272:408-411; Yang et al., 1996, Proc. Natl. Acad. Sci. USA 93:12845-12850; Emiliani et al., 1998, Proc. Natl. Acad. Sci. USA 95:2795-10 2800), and yeast RPD3 (Videl & Gaber, 1991, Mol. Cell. Biol. 11:6317-6327), and class II by the human HDAC4, 5, 6 (Grozinger et al., 1999, Proc. Natl. Acad. Sci. USA 96:4868-4873; Fischle, et al., 1999, J. Biol. Chem. 274:11713-15 11720), and yeast HDA1 (Rundlett et al., 1996, Proc. Natl. Acad. Sci. USA 93:14503-14508). The two classes share a ~390 amino acid region of sequence similarity, comprising the deacetylase core, but are divergent outside this region. The histone deacetylase genes belong to an even larger 20 superfamily (Leipe & Landsman, 1997, Nucleic Acids Res. 25:3693-3697) that contains the prokaryotic utilization proteins (AcuC; 28.1% sequence identity to HDAC1), and the prokaryotic acetylpolyamine amidohydrolases (APAH; 15.0 % sequence identity to HDAC1). The enzymatic activity of AcuC is not clear, but its disruption reduces 25 the ability of B. subtilis to breakdown acetoin and utilize it as a carbon source. Grundy et al., 1993, Mol. Microbiol. 10:259-271. APAHs catalyze the deacetylation of polyamines by cleaving a non-peptide amide bond (reviewed in Leipe & Landsman, 1997, Nucleic Acids Res. 25:3693-3697). 30

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It is useful to address the questions of how HDACs and HDACrelated proteins catalyze the deacetylation of histones and how the above-referenced compounds, particularly those compounds with antitumor activity, inhibit this activity in order to better understand the mechanism of inhibition of HDACs and to facilitate discovery of additional useful compounds which may inhibit this activity. To this end, the present invention has determined the three dimensional structure of a HDAC1-like protein from the thermophilic bacterium Aquifex aeolicus, herein after HDLP. determination of the nucleic acid coding sequence of HDLP was described by Deckert et al., 1998, Nature 392:353-358. encoded 375 residue protein, whose sequence was The determined from the nucleic acid encoding sequence, shares 35.2% amino acid sequence identity with HDAC1, deacetylates histones in vitro, and is inhibited by TSA, SAHA and several other HDAC inhibitors. The determination of the threedimensional structure of HDLP is useful in the design, identification and screening of new HDAC family inhibitory compounds which are useful for the inhibition of cell growth both in vivo and in vitro.

Summary of the Invention

In general, it is the object of the present invention to provide detailed three-dimensional structural information for a family of proteins known as histone deacetylases (HDAC), and particularly a homologue from the hyperthermophilic bacterium Aquifex aeolicus HDLP (histone deacetylase-like protein) which shares 35.2 % sequence identity with human histone deacetylase (HDAC1). It is also an object of the present invention to provide three-

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dimensional structural information of an HDLP bound to an inhibitory compound.

In one embodiment of the invention, three-dimensional structure information is obtained from a crystal of wildtype HDLP (SEQ ID NO:1) (the nucleic acid encoding wild-type In a further embodiment of the HDLP is SEQ ID NO:2). invention, three-dimensional information is obtained from a mutant HDLP comprising two mutations (1) cysteine 75 to a serine and (2) cysteine 77 to a serine (Cys75Ser/Cys77Ser double mutant; SEQ ID NO:3) (the nucleic acid encoding HDLP Cys75Ser/Cys77Ser double mutant is SEQ ID NO:4). The HDLP ofthe invention facilitates mutant present the determination of three-dimensional structural information of HDLP bound to a zinc atom at its zinc atom-binding site.

In a preferred embodiment of the invention, the threedimensional structural information is obtained from a cocrystal of a protein-inhibitor compound complex that comprises HDLP or HDLP Cys75Ser/Cys77Ser double mutant and trichostatin A (TSA). In another preferred embodiment of the invention the three-dimensional structural information is obtained from a co-crystal of a protein-inhibitor compound complex that comprises HDLP or HDLP Cys75Ser/Cys77Ser double suberoylanilide mutant and hydroxamic acid (SAHA). Any HDLP or HDLP-related protein (e.g. HDAC) inhibitor compound that may be co-crystallized with HDLP may be used to form a co-crystal of the present invention.

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The protein crystals and protein-inhibitory complex cocrystals of the present invention diffract to a high 5

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resolution limit of at least equal to or greater than 4 angstrom (Å). In a preferred embodiment, the protein crystals and protein-inhibitory complex co-crystals of the present invention diffract to a high resolution limit of greater than 2.5 Å.

A crystal of the present invention may take a variety of forms, all of which are contemplated by the present invention. In a preferred embodiment, the crystal has a space group of C2 with one molecule in the asymmetric unit and with unit dimensions of a = 51.4 Å, b = 93.8 Å, 78.7 Å and β = 96.9° (see, e.g., Example 2, below). another preferred embodiment, the crystal has a space group of $P2_12_12_1$ with two molecules in the asymmetric unit and with unit dimensions of a = 53.4 Å, b = 94.4 Å, c = 156.3 Å (see, e.g., Example 2, below). The HDLP structure comprises a parallel β sheet with α helices packing against both faces. At one end of the β sheet, the HDLP has a narrow, tube-like pocket formed by several well-ordered loops. The walls of the pocket are lined with hydrophobic residues and there is a zinc binding site and several polar side chains at the bottom of the pocket. The inhibitory compounds of the present invention bind in the pocket.

The three-dimensional structural information obtained from crystals of HDLP, HDLP Cys75Ser/Cys77Ser double mutant, HDLP Cys75Ser/Cys77Ser double mutant comprising a zinc atom, HDLP comprising an inhibitory compound such as TSA or SAHA, and HDLP Cys75Ser/Cys77Ser double mutant comprising an inhibitor compound such as TSA or SAHA may be employed to solve the structure of any HDLP-related protein (e.g. HDAC) crystal,

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or any mutant HDLP-related protein and particularly any wild type or mutant of HDLP-related protein complexed with a ligand, including a substrate or inhibitor compound. If the crystals are in a different space group than the known structure, molecular replacement may be employed to solve the structure, or if the crystals are in the same space group, refinement and difference fourier methods may be employed. The structure of HDLP-related proteins (e.g. HDAC1) comprise no greater than a 2.0 Å root mean square deviation (rmsd) in the positions of the Ca atoms for at least 50% or more of the amino acids of the full-length HDLP structure.

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The present invention also provides a nucleic acid molecule encoding an HDLP Cys75Ser/Cys77Ser double mutant having the amino acid sequence of SEQ ID NO:3 and the nucleic acid sequence of SEQ ID NO:4. It is also contemplated by the invention that mutations be made in HDLP-related proteins at cysteine residues, as with the Cys75Ser/Cys77Ser double mutant, in order to facilitate the determination of the structure of said proteins bound to a zinc atom. Additionally, the present invention provides expression vectors which comprise the nucleic acid molecule encoding an HDLP Cys75Ser/Cys77Ser double mutant encoded by the sequence represented by SEQ ID NO:4 operatively linked to expression control sequences.

It is another object of the present invention to provide methods for the design, identification and screening of potential inhibitor compounds of the HDLP/HDAC family. In a preferred embodiment the method for the rational design,

identification and screening of potential inhibitor compounds for HDLP and HDLP-related proteins (e.g. HDACs) comprising deacetylase activity comprises the steps of: (a) using a three-dimensional structure of an HDLP as defined by the atomic coordinates of the present invention; employing said three-dimensional structure to design or select said potential inhibitor compound; (c) synthesizing and/or selecting said potential inhibitor; (d) contacting said potential inhibitor compound with said enzyme in the presence of acetylated substrate; and (e) determining the percent inhibition of deacetylase activity to determine the inhibitory activity of said potential inhibitor compound. In a further preferred embodiment, the binding properties of said rationally designed inhibitory compound may be determined by a method comprising the steps of: (a) forming a complex comprising said inhibitory compound and HDLP or a HDLP-related protein, (b) co-crystallizing said inhibitory compound-HDLP complex; (c) determining said dimensional structure of said co-crystal through molecular replacement or refinement and difference fourier with the molecular coordinates of HDLP as defined by the present invention; and (d) analyzing the three-dimensional structure to determine the binding characteristics of said potential inhibitor compound.

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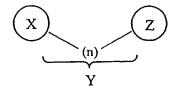
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It is a further object of the present invention to identify a defined class of HDLP/HDAC family inhibitor compounds. The HDLP/HDAC family inhibitor compounds of the present invention are represented by formula (I):

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(I)



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wherein X comprises a cap group which binds to at least one amino acid selected from the group consisting of proline and leucine; Y comprises an aliphatic chain group which binds to at least one amino acid selected from the group consisting of leucine, phenylalanine and glycine; and Z comprises and active site binding group which binds to at least one amino acid selected from the group consisting of aspartic acid, tyrosine and histidine and may further bind to a zinc atom.

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Brief Description of the Drawings

Figure 1 is a table listing the statistics from the X-ray crystallographic analysis of a HDLP crystal, a HDLP-TSA co-crystal, and a HDLP-SAHA co-crystal.

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Figure 2 shows an alignment of various HDAC homologues with percent sequence identity depicted.

Figure 3 shows a graph indicating the histone deacetylase activity of HDLP and HDAC1 and the inhibition of HDLP and HDAC1 by the inhibitors TSA and HC-toxin.

Figure 4 shows (A & B) a schematic representation of the HDLP-Zn²⁺-TSA complex in two approximately orthogonal views,

(C) a topology diagram of HDLP indicating the regions of homology with HDAC1, and (D) a close-up schematic representation of the HDLP-Zn²⁺-SAHA complex.

Figure 5 shows (A) a schematic representation of a slice through a surface representation of HDLP with the pocket internal cavities and position of the β sheet indicated, (B) a schematic representation of a close-up view of the active site looking down into the pocket in an orientation similar to Figure 4B.

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Figure 6 shows (A) a space-filling representation of TSA in the active site pocket, (B) a closeup stereo view of the structure of the HDLP-ZN²⁺-TSA complex in a similar orientation to Figure 4B, and (C) a schematic representation of the HDLP-TSA interactions.

Figure 7 shows (A) a schematic representation of the regions of homology shared between HDLP and HDAC1 in an orientation similar to that of Figure 4A, and (B) a detailed schematic representation of the homology shared in the pocket and internal cavity between HDLP and HDAC1 in an orientation similar to that of Figure 4B.

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Figure 8 shows a schematic representation of the proposed catalytic mechanism for the deacetylation of acetylated lysine.

Figure 9 shows a schematic representation of a space filling diagram showing the conserved amino acids in the active site and nearby grooves.

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Figure 10 is the nucleic acid sequence of HDLP from Aquifex aeolicus (SEQ ID NO. 2).

Figure 11 is the amino acid sequence of full length HDLP from Aquifex aeolicus (SEQ ID NO. 1).

Figure 12 is the nucleic acid sequence of the HDLP active site mutant Tyr297Phe (SEQ ID NO. 6).

Figure 13 is the amino acid sequence of the HDLP active site mutant Tyr297Phe (SEQ ID NO. 5).

Figure 14 is the nucleic acid sequence of a double mutant of HDLP from Aquifex aeolicus comprising a Cys75Ser and Cys77Ser mutation (SEQ ID NO. 4).

Figure 15 is the amino acid sequence of a double mutant of HDLP from Aquifex aeolicus comprising a Cys75Ser and Cys77Ser mutation (SEQ ID NO. 3).

- Figure 16-1 to 16-49 lists the atomic structure coordinates for HDLP as derived by X-ray diffraction from a crystal of HDLP.
- Figure 17-1 to 17-49 lists the atomic structure coordinates

 for HDLP Cys75Ser/Cys77Ser double mutant comprising a zinc
 atom in the active site as derived by X-ray diffraction from
 a crystal of the HDLP Cys75Ser/Cys77Ser double mutant.
- Figure 18-1 to 18-99 lists the atomic structure coordinates

 for HDLP Cys75Ser/Cys77Ser double mutant as derived by X-ray
 diffraction from a co-crystal of HDLP complexed with TSA.
- Figure 19-1 to 19-48 lists the atomic structure coordinates for HDLP Cys75Ser/Cys77Ser double mutant as derived by X-ray diffraction from a co-crystal of HDLP complexed with SAHA.

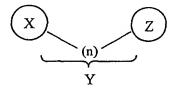
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Detailed Description of the Invention

The present invention provides crystals of a histone deacetylase (HDAC) homologue grown in the presence and absence of a compound capable of inhibiting the histone deacetylase activity of said HDAC homologue. As referred to herein, a HDAC homologue (as well as a HDLP-related protein) is any protein molecule having (a) greater than 15% sequence identity to over the 375 amino acid residues of HDLP; (b) having no more than twenty insertions or deletions for a total of no more than 100 amino acids; and (c) deacetylase activity. Sequence identity is calculated by the program DNAstar™ using the identity matrix weighing scheme clustal method (DNAstar program, Madison, WI).

A HDLP/HDAC inhibitor compound, as used herein, refers to any compound represented by Formula (I):

(I)



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wherein X comprises a cap group which binds to at least one amino acid selected from the group consisting of tyrosine, proline and leucine; Y comprises an aliphatic chain group from about 5 to about 10 Å, preferably 7Å, which binds to at least one amino acid selected from the group consisting of phenylalanine and glycine; and Z comprises a active site binding group which binds to at least one amino acid selected from the group consisting of aspartic acid, tyrosine and histidine and which may further bind to a zinc atom. The HDAC inhibitory compounds of the present

invention can inhibit greater than 50% of the histone deacetylase activity of a HDAC homologue or a HDLP-related protein.

To grow the crystals of the present invention, the HDAC and HDAC-inhibitory compound complex are purified to greater than 80% total protein and more preferably purified to greater than 90% total protein. For expression and purification purposes, the full-length HDLP (Genbank accession number AE000719) may be subcloned from Aquifex aeolicus chromosomal DNA preparation by the polymerase chain reaction (PCR) and inserted into an expression vector.

A large number of vector-host systems known in the art may be used. Possible vectors include, but are not limited to, 15 plasmids or modified viruses, but the vector system must be compatible with the host cell used. Examples of vectors include E. coli bacteriophages such as lambda derivatives, or plasmids such as pBR322 derivatives or pUC plasmid 20 derivatives, e.g., pGEX vectors (Amersham-Pharmacia, Piscataway, New Jersey), pET vectors (Novagen, Madison, WI), pmal-c vectors (Amersham-Pharmacia, Piscataway, New Jersey), pFLAG vectors (Chiang and Roeder, 1993, Pept. Res. 6:62-64), baculovirus vectors (Invitrogen, Carlsbad, CA; Pharmingen, San Diego, CA), etc. 25 The insertion into a cloning vector can, for example, be accomplished by ligating the DNA fragment into a cloning vector which has complementary cohesive termini, by blunt end ligation if no complementary cohesive termini are available or by through nucleotide linkers using techniques standard in the art. 30 Ausubel et al. (eds.), Current Protocols in Molecular

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Biology, (1992). Recombinant vectors comprising the nucleic acid of interest may then be introduced into a host cell compatible with the vector (e.g. E. coli, insect cells, mammalian cells, etc.) via transformation, transfection, infection, electroporation, etc. The nucleic acid may also be placed in a shuttle vector which may be cloned and propagated to large quantities in bacteria and then introduced into a eukaryotic cell host for expression. The vector systems of the present invention may provide expression control sequences and may allow for the expression of proteins in vitro.

In a preferred embodiment, the full length HDLP (SEQ ID NO:2) is subcloned from Aquifex aeolicus chromosomal DNA preparation into pGEX4T3 (Amersham-Pharmacia, Piscataway, New Jersey). In order to construct a double mutant comprising a Cys75Ser and Cys77Ser mutation (SEQ ID NO:4), and to construct the HDLP active site mutant Tyr297Phe (SEQ ID NO:5 and SEQ ID NO:6), PCR site directed mutagenesis may be employed with verification by DNA sequencing by methods known to those skilled in the art (see, e.g., Exemple 1 below). The mutants of the present invention may be subcloned into a suitable expression vector and introduced into a host cell for protein production, as described above.

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The HDLP nucleic acids of the present invention may be subcloned into an expression vector to create an expression construct such that the resultant HDLP molecule which is produced comprises a fusion protein wherein said fusion protein comprises a tag for ease of purification. As referred to herein, a "tag" is any additional amino acids

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which are provided in a protein either c-terminally, nterminally or internally for the ease of purification, for the improvement of production or for any other purpose which may facilitate the goals of the present invention (e.g. to achieve higher levels of production and/or purification). Such tags include tags known to those skilled in the art to be useful in purification such as, but not limited to, his tag, glutathione-s-transferase tag, flag tag, mbp (maltose binding protein) tag, etc. In a preferred embodiment, the wild-type and mutant HDLPs of the present invention are tagged with glutathione-s-transferase (see Example 1 below). In another preferred embodiment, HDAC1 is flag tagged (see Example 1 below). Such tagged proteins may also be engineered to comprise a cleavage site, such as a thrombin, enterokinase or factor X cleavage site, for ease of removal of the tag before, during or after purification. systems which provide a tag and a cleavage site for removal of the tag are particularly useful to make the expression constructs of the present invention.

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The tagged HDLPs and HDACs of the present invention may be purified by immuno-affinity or conventional chromatography, including but not limited to, chromatography employing the following: glutathione-sepharose™ (Amersham-Pharmacia, Piscataway, New Jersey) or an equivalent resin, nickel or cobalt-purification resins, anion exchange chromatography, cation exchange chromatography, hydrophobic resins, gel filtration. antiflag epitope resin, reverse chromatography, etc. After purification, the HDLP and HDLPinhibitor compound complex may be concentrated to greater than 1 mg/ml for crystallization purposes. In a preferred embodiment HDLP and HDLP-inhibitor complexes

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concentrated to greater than 10 mg/ml for crystallization and in a particularly preferred embodiment, HDLP and HDLP-inhibitor complexes are concentrated to greater than 20 mg/ml.

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In order to determine whether the purified HDLPs of the present invention demonstrate histone deacetylase activity, the purified HDLPs and also any HDLP-related protein may be assayed by any method known to those skilled in the art for the determination of said activity. In a preferred embodiment, the purified HDLPs of the present invention are incubated in the presence of [3H] acetyl-labeled histone substrate (Carmen et al., 1996, J. Biol. Chem. 271:15837-15844) in a buffer suitable for detection of histone deacetylase activity (see Example 3 below); stopping the reaction; extracting the released acetate and measuring said released acetate, as described by Henzel et al. (J. Biol. Chem. 266:21936-21942 (1991); Example 3 below). preferred embodiment, the HDLPs of the present invention are inclubated in the presence of ZnCl2 in order to obtain histone deacetylase activity therefrom (Example 3 below).

In another embodiment, the crystals of the present invention comprise purified wild-type HDLP (SEQ ID NO:1) and are grown at room temperature by the hanging-drop vapor-diffusion method from a crystallization solution comprising one or more precipitants selected from the group consisting of isopropanol, polyethylene glycol, and tert butanol (see Example 2 below). The crystallization solution may further comprise one or more salts including salts selected from the group consisting of NaCl and KCl, and one or more buffers

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including buffers selected from the group consisting of Tris (tris(hydroxymethyl)aminomethane and bis-tris propane-Cl (1,3-bis[tris(hydroxymethyl)methyl-amino] propane) (see Example 2 below). The pH of the crystallization solution is preferably between pH 5 to 9, although other pH values are also contemplated by the present invention (see Example 2 below).

Any crystallization technique known to those skilled in the art may be employed to obtain the crystals of the present invention, including, but not limited to, batch crystallization, vapor diffusion (either by sitting drop or hanging drop) and micro dialysis. Seeding of the crystals in some instances may be required to obtain X-ray quality crystals. Standard micro and/or macro seeding of crystals may therefore be used.

The crystals of the present invention may form in the space group C2 with one molecule in the asymmetric unit and with unit dimensions of a=51.4 Å, b=93.8 Å, c=78.7 Å and $\beta=96.9^{\circ}$ (see Example 2 below). The crystals of the present invention may also form in the space group $P2_12_12_1$ with two molecules in the asymmetric unit and with unit dimensions of a=53.4 Å, b=94.4 Å, c=156.3 Å (see Example 2 below). However, the present invention contemplates crystals which form in any space group including, but not limited to, C2, $P2_1$, $P2_12_12_1$, $P3_121$, $P4_22_12_1$, and $C222_1$. The crystals diffract to a resolution greater than 4 Å, preferably greater than 2.5 Å.

To collect diffraction data from the crystals of the present

invention, the crystals may be flash-frozen in crystallization buffer employed for the growth of said crystals, however with preferably higher precipitant concentration (see, e.g., Example 2 below). For example, but not by way of limitation, if the precipitant used was 28% PEG 1500, the crystals may be flash frozen in the same crystallization solution employed for said crystal growth wherein the concentration of the precipitant is increased to 35% (see Example 2 below). If the precipitant is not a sufficient cryoprotectant (i.e. a glass is not formed upon cryoprotectants (e.g. glycerol, low flash-freezing), molecular weight PEGs, alcohols, etc) may be added to the solution in order to achieve glass formation upon flashfreezing, providing the cryoprotectant is compatible with preserving the integrity of the crystals. The flash-frozen crystals are maintained at a temperature of less than -110°C and preferably less than -150°C during the collection of the crystallographic data by X-ray diffraction. diffraction data may be processed with DENZO and SCALEPACK (Otwinowski & Minor, 1997, Method Ensemble. 276:307-326) but any method known to those skilled in the art may be used to process the X-ray diffraction data.

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In order to determine the atomic structure of HDLP according to the present invention, multiple isomorphous replacement (MIR) analysis, model building and refinement may be performed. For MIR analysis, the crystals may be soaked in heavy-atoms to produce heavy atom derivatives necessary for MIR analysis. As used herein, heavy atom derivative or derivitization refers to the method of producing a chemically modified form of a protein or protein complex

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crystal wherein said protein is specifically bound to a heavy atom within the crystal. In practice a crystal is soaked in a solution containing heavy metal atoms or salts, or organometallic compounds, e.g., lead chloride, gold cyanide, thimerosal, lead acetate, uranyl acetate, mercury chloride, gold chloride, etc, which can diffuse through the crystal and bind specifically to the protein. location(s) of the bound heavy metal atom(s) or salts can be determined by X-ray diffraction analysis of the soaked This information is used to generate MIR phase crystal. information which is used to construct the three-dimensional structure of the crystallized HDLPs and HDLP-related proteins of the present invention. In a preferred embodiment, the heavy atoms comprise thimerosal, $\text{KAu}\left(\text{CN}\right)_2$ and $Pb(Me)_3OAc$ (see Example 2 below). The MIR phases may be calculated by any program known to those skilled in the art and preferably with the program MLPHARE (The CCP4 suite: Programs for computational crystallography, 1994, Crystallogr. D. 50:760-763) and may also use the anomalous diffraction signal from the thimerosal derivative. preferred embodiment, the MIR phases were calculated at 2.5 Å and have a mean figure of merit of 0.55 (see Figure 19 and Example 2 below). The phases may be improved where necessary by solvent flattening by methods known to those skilled in the art including, but not limited to, through the use of the program DM (The CCP4 suite: Programs for computational crystallography, 1994, Acta Crystallogr. D <u>50</u>:760-763).

30 Thereafter, an initial model of the three-dimensional structure may be built using the program O (Jones et al.,

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1991, Acta Crystallogr. A <u>47</u>:110-119). The interpretation and building of the structure may be further facilitated by use of the program CNS (Brunger et al., 1998, Acta Crystallogr. D <u>54</u>:905-921).

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For the determination of the HDLP-inhibitor compound complex structure, if the space group of the HDLP-inhibitor compound complex crystal is different, molecular replacement may be employed using a known structure of apo-HDLP (as referred to herein, apo-HDLP or apo-HDAC is the enzyme which is not complexed with an inhibitor compound) or any known HDLP/inhibitor complex structure whose structure may be determined as described above and below in Example 2. Ιf the space group of the HDLP-inhibitor compound crystals is the same, then rigid body refinement and difference fourier may be employed to solve the structure using a known structure of apo-HDLP (as referred to herein, apo-HDLP or apo-HDAC is the enzyme which is not complexed with an inhibitor compound) or any known HDLP/inhibitor complex structure.

The term "molecular replacement" refers to a method that involves generating a preliminary model of the three-dimensional structure of the HDLP crystals of the present invention whose structure coordinates are unknown prior to the employment of molecular replacement. Molecular replacement is achieved by orienting and positioning a molecule whose structure coordinates are known (in this case the previously determined apo-HDLP) within the unit cell as defined by the X-ray diffraction pattern obtained from an HDLP or HDLP-related protein crystal whose structure is

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unknown so as to best account for the observed diffraction pattern of the unknown crystal. Phases can then be calculated from this model and combined with the observed amplitudes to give an approximate Fourier synthesis of the structure whose coordinates are unknown. This in turn can be subject to any of several forms of refinement to provide a final, accurate structure.

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Any method known to the skilled artisan may be employed to 10 determine the structure by molecular replacement. For example, the program AMORE (The CCP4 suite: Programs for computational crystallography, 1994, Acta Crystallogr. D. 50:760-763) may be employed to determine the structure of an unknown histone deacetylase +/- an inhibitor by molecular replacement using the apo-HDLP coordinates (Figure 16). For the structure determination of the inhibitory compound TSA, the structure of TSA was obtained from the Cambridge Structural Database (Refcode TRCHST, http://www.ccdc.cam.ac.uk >>) may be employed to define the stereochemical restraints used in the refinement with the program CNS (Brunger et al., 1998, Acta Crystallogr. D <u>54</u>:905-921).

The three-dimensional structural information and the atomic coordinates associated with said structural information of HDLP are useful for solving the structure of crystallized proteins which belong to the HDAC family by molecular Similarly, any structure of a crystallized replacement. protein which is thought to be similar in structure based on function or sequence similarity or identity to HDLP may be solved by molecular replacement with the HDLP structural

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information of the present invention. The structure of HDLP-related proteins as determined by molecular replacement as described above and in Example 2 below, comprise a root mean square deviation (rmsd) of no greater than 2.0 Å in the positions of $C\alpha$ atoms for at least 50% or more of the amino acids of the structure over the 375 residues of full-length HDLP. Such a rmsd may be expected based on the amino acid sequence identity. Chothia & Lesk, 1986, Embo J. $\underline{5}$:823-826.

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The refined three-dimensional HDLP structures of the present invention, specifically apo-HDLP, Cys75Ser/Cys77Ser double mutant HDLP comprising a zinc atom in the active site, HDLP/TSA complex comprising a zinc atom in the active site, and HDLP/SAHA complex comprising a zinc atom in the active site, are represented by the atomic coordinates set forth in Figures 16 to 19 respectively. The refined model for apo-HDLP comprising amino acids 1-375 consists of wild-type HDLP residues 2 to 373 with residues 1, 374 and 375 not modeled and presumed disordered and was determined to a resolution Â. 1.8 Similarly, the refined model Cys75Ser/Cys77Ser double mutant HDLP comprising a zinc atom in the active site also consists of residues 2 to 373 with residues 1, 374 and 375 not modeled and presumed disordered and was determined to a resolution of 2.0 Å. The refined model for the HDLP/TSA complex comprising a zinc atom in the active site consists of the Cys75Ser/Cys77Ser double mutant HDLP residues 2 to 373 with residues 1, 374 and 375 not modeled and presumed disordered, has TSA in the binding pocket and was determined to a resolution of 2.1 Å. HDLP/SAHA complex is similar to the HDLP/TSA complex but has SAHA in the binding pocket and was determined to a resolution of 2.5 Å.

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For the purposes of further describing the structure of HDLP and HDLP-related proteins, including, but not limited to, HDACs, from the data obtained from the HDLP crystals of the present invention, the definition of the following terms is provided:

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The term " β sheet" refers to two or more polypeptide chains (or β strands) that run alongside each other and are linked in a regular manner by hydrogen bonds between the main chain C=O and N-H groups. Therefore all hydrogen bonds in a beta-sheet are between different segments of polypeptide. Most β -sheets in proteins are all-parallel (protein interiors) or all-antiparallel (one side facing solvent, the other facing the hydrophobic core). Hydrogen bonds in antiparallel sheets are perpendicular to the chain direction and spaced evenly as pairs between strands. Hydrogen bonds in parallel sheets are slanted with respect to the chain direction and spaced evenly between strands.

The term " α helix" refers to the most abundant helical 20 conformation found in globular proteins. The average length of an α helix is 10 residues. In an α helix, all amide protons point toward the N-terminus and all carbonyl oxygens point toward the C-terminus. The repeating nature of the phi, psi pairs ensure this orientation. Hydrogen bonds 25 within an $\boldsymbol{\alpha}$ helix also display a repeating pattern in which the backbone C=O of residue X (wherein X refers to any amino acid) hydrogen bonds to the backbone HN of residue X+4. The α helix is a coiled structure characterized by 3.6 residues per turn, and translating along its axis 1.5 Å per 30 amino acid. Thus the pitch is 3.6x1.5 or 5.4 Å. The screw sense of alpha helices is always right-handed.

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The term "loop" refers to any other conformation of amino acids (i.e. not a helix, strand or sheet). Additionally, a loop may contain bond interactions between amino acid side chains, but not in a repetitive, regular fashion.

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Amino acid residues in peptides shall herein after be abbreviated as follows: Phenylalanine is Phe or F; Leucine is Leu or L; Isoleucine is Ile or I; Methionine is Met or M; Valine is Val or V; Serine is Ser or S; Proline is Pro or P; Threonine is Thr or T; Alanine is Ala or A; Tyrosine is Tyr or Y; Histidine is His or H; Glutamine is Gln or Q; Asparagine is Asn or N; Lysine is Lys or K; Aspartic Acid is Asp or D; Glutamic Acid is Glu or E; Cysteine is Cys or C; Tryptophan is Trp or W; Arginine is Arg or R; and Glycine is Gly or G. For further description of amino acids, please refer to Proteins: Structure and Molecular Properties by Creighton, T.E., W.H. Freeman & Co., New York 1983.

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The term "positively charged amino acid" refers to any amino acid having a positively charged side chain under normal physiological conditions. Examples of positively charged amino acids are Arg, Lys and His. The term "negatively charged amino acid" refers to any amino acid having a negatively charged side chain under normal physiological conditions. Examples of negatively charged amino acids are Asp and Glu. The term "hydrophobic amino acid" refers to any amino acid having an uncharged, nonpolar side chain that is relatively insoluble in water. Examples of hydrophobic amino acids are Ala, Leu, Ile, Gly, Val, Pro, Phe, Trp and Met. The term "hydrophilic amino acid" refers to any amino acid having an uncharged, polar side chain that is

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relatively soluble in water. Examples of hydrophilic amino acids are Ser, Thr, Tyr, Asp, Gln, and Cys. The term "aromatic amino acid" refers to any amino acid comprising a ring structure. Examples of aromatic amino acids are His, Phe, Trp and Tyr.

The term "charge relay system" refers to a His-Asp arrangement as described by Fersht & Sperling, 1973, J. Mol. Biol. 74:137-149; Blow et al., 1969, Nature 221:337-340.

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The information obtained from the three-dimensional structures of the present invention reveal that HDLP has a single-domain structure that belongs to the open α/β class of folds (see, e.g., Branden, 1980, Q. Rev. Biophys. 13:317-38). Two orthogonal views of the overall three-15 dimensional structure of HDLP are depicted in Figure 4A and The HDLP structure has a central eight-stranded 4B. parallel β sheet (strands arranged as $\beta2\text{-}\beta1\text{-}\beta3\text{-}\beta8\text{-}\beta7\text{-}\beta4\text{-}\beta5\text{-}$ $\beta6)$, and sixteen α helices (labeled $\alpha1$ through $\alpha16$ respectively). See Figure 4C. Four of the helices pack on 20 either face of the β sheet ($\alpha 7,~\alpha 8,~\alpha 9,~\alpha 10$ and $\alpha 11,~\alpha 12,$ α 13, α 14) forming the core α/β structure characteristic of this class of folds. Most of the remaining eight helices are positioned near one side of the β sheet, near stands $\beta 2$ - β 1- β 3- β 8. Large, well defined loops (Loops L1-L7; Figure 25 4C) originate from the C-terminal ends of the β -strands. The extra helices and the large L1-L7 loops are associated with a significant extension of the structure beyond the core α/β motif. This extension of the structure gives rise to two prominent architectural features: a deep, narrow 30 pocket and an internal cavity adjacent to the pocket. These

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two architectural features comprise the active site (see Figure 5A). The structure of HDLP-related proteins (e.g. HDACs) may also comprise the conserved α/β structure characteristic.

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The term "active site" comprises any or all of the following sites in HDLP, the substrate binding site, the site where the cleavage of an acetyl group from a substrate occurs or the site where an inhibitor of the HDAC family or, more particularly, HDLP binds. The active site, as referred to herein, comprises Aspl66, Asp258, His170, Tyr297, His131, His132, Asp168, Asp173, Phe141, Phe198, Leu265, Pro22 and Gly140, and also a metal bound at the bottom of the pocket by Asp173, Asp168 and His defined by the coordinates listed in Figures 16 to 19 with an rmsd of 2.0 Å. The metal which binds at the bottom of the pocket will be a divalent cation selected from the group consisting of zinc, cobalt or manganese.

The deep narrow pocket has a tube-like shape with a depth of ~ 11 Å. The pocket opening constricts half way down to ~ 4.5 by 5.5 Å, and becomes wider at the bottom (see Figure 5A). The pocket and its immediate surroundings are made up of loops L1 through L7.

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The walls of the pocket are covered with side chains of hydrophobic and aromatic residues (Pro22, Tyr91 near the entrance; and Gly140, Phe141, Phe 198, Leu265 and Tyr297 further down; Figure 5B). For numbering of amino acids please refer to SEQ ID NO:1. Of particular interest are Phe141 and Phe198, whose phenyl groups face each other in

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parallel at a distance of 7.5 Å, marking the most slender portion of the pocket (see Figure 5B). Of particular interest is that only one pocket residue differs in HDAC1 when the sequences are aligned (alignment may be accomplished using DNAstar™ MegAlign™ program, Madison, WI), this residue is Glu98 of HDAC1 which is Tyr91 in HDLP. The structure reveals that this residue in HDLP is mostly solvent exposed.

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Near the bottom of the pocket of the active site at its narrowest point, is located a zinc ion (see Figure 6A). In order to obtain the zinc in the structure, the crystals may be soaked in zinc (e.g. ZnCl₂) or co-crystalized in the presence of zinc. The zinc ion is coordinated by Aspl68 (0δ1, 2.1 Å), His170 (Nδ1, 2.1 Å), Asp258 (0δ1, 1.9 Å) and a water molecule (2.5 Å). See Figure 5B and 6B. The amino acid residues that coordinate zinc are arranged in a tetrahedral geometry, but the position of the water molecule, which is also hydrogen bonded to His131, deviates from this geometry by ~25°.

In addition to the zinc ligands, the bottom of the pocket contains two histidine (His131 and His132), two aspartic acids (Asp166 and Asp173) and a tyrosine (Tyr297). See Figure 5B and 10B. Each of the histidines makes a hydrogen bond through its N δ 1 to an aspartic acid carboxylate oxygen, with the oxygen located in the plane of the imidizole ring (Figure 5B). This His-Asp arrangement is characteristic of the charge relay system present in the active sites of serine proteases, where it serves to polarize the imidizole Ne and increase its basicity. Fersht & Sperling, 1973, J.

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Mol. Biol. <u>74</u>:137-149; Blow et al., 1969, Nature <u>221</u>:337-340.

The Asp166-His131 charge pair relay (hereafter referred to as "buried charged relay") is positioned even deeper in the pocket and more buried compared to the Asp173-His132 charge relay (hereafter referred to as "exposed charge relay") which is partially solvent exposed. The buried charge relay makes a hydrogen bond (2.6 Å) to the zinc-bound water molecule referred to above, and this hydrogen bond could contribute to the deviation of the water-zinc coordination from ideal geometry (Figure 5B). The exposed charge relay is directed to a point ~ 2.5 Å away from the water molecule and closer to the surface.

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Tyr 297 is positioned next to the zinc, opposite from where the two charge relay systems are located. The Tyr hydroxyl group lies 4.4 Å away from the zinc atom and has no interactions with the rest of the protein (Figure 5B). Next to Tyr297, there is an opening in the pocket wall, which leads to the adjacent internal cavity.

The floor of the internal cavity is made up of portions of the L3 and L7 loops as they emerge from the β strands, and the roof is made up by the $\alpha 1\text{-L}1\text{-}\alpha 2$ segment. The L1 loop appears more flexible than other loops in the structure. This may allow the transient exchange of the cavity contents with the bulk solvent.

The cavity is lined primarily with hydrophobic residues and is particularly rich in glycine residues (Ala127, Gly128, Gly129, Met130, and Phe141 of L3; Gly293, Gly294, Gly295 and

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Gly296 of L7; and Tyr17, Pro22 and Leu23 of L1). There are only two charged residues in the cavity (Arg27 and His 21) and these are contributed by the L1 loop.

The cavity may provide space for the diffusion of the acetate product away from the catalytic center, which may otherwise be crowded and shielded during deacetylation from the solvent when the substrate is bound. Such a role for the cavity is supported by the observation that the cavity contains three water and two isopropanol molecules (from the crystallization buffer) in the 1.8 Å apo-protein structure. The cavity may also bind another cofactor, in addition to zinc, for the facilitation of the enzymatic activity of the HDLP. A proposed catalytic mechanism for deacetylation is provided in Figure 8.

The structure of HDLP as defined by the present invention, in conjunction with the HDAC1 sequence homology, shows that the 375-amino acid HDLP protein corresponds to the histone deacetylase catalytic core which is conserved across the HDAC family (see Figure 2). The 35.2% HDLP-HDAC1 sequence identity predicts structural similarity with a rmsd in $\ensuremath{\text{C}\alpha}$ positions of ~ 1.5 Å. Chothia and Lesk describe the relation between the divergence of sequence and structure of proteins in Embo J. $\underline{5}$:823-826 (1986). residue C-terminus of HDLP is likely to have a divergent structure since this region has lower homology to HDAC1, although the $\alpha 16$ helix in this region is part of the conserved open α/β core fold and HDAC1 is likely to comprise a similar helix. However divergent this C-terminal region may be, this region is outside the active site and is likely to not effect the structure of the active site. Beyond the

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C-terminus of the histone deacetylase catalytic core, HDAC family members are divergent in length and sequence. In the HDAC family, this region (amino acid residues ~390-482) is highly polar, populated with acidic residues, and is likely to be flexible or loosely folded.

The HDLP-HDAC homology maps primarily to the hydrophobic core and to the L1-L7 loops, with portions of the loops that make up the pocket and adjacent cavity having the highest level of amino acid residue sequence conservation (Figure 9A and 9B). Specifically, all of the polar residues in the active site (the zinc ligands, the two charge relay systems, and Tyr297) and the hydrophobic residues that make up the walls of the pocket (Gly140, Phe141, Phe198 and Leu265) are identical. Among the residues that make up the internal cavity, the ones closest to the active site are either identical or conservatively substituted (for example, Leu23 \rightarrow Met and Met130 \rightarrow Leu). Surface residues around the pocket are conserved to a lesser extent, but are still above 35% average sequence identity.

The information obtained from the inhibitor-bound HDLP complex crystal structures of the present invention reveal detailed information which is useful in the design, isolation, screening and determination of potential inhibitor compounds which may inhibit HDLP/HDAC family members. As described above, the HDLP structure consists of a parallel β sheet with α helices packing against both faces (Figure 4A, 4B, and 4C). At one end of the β sheet, 7 loops (L1-L7) form a narrow, tube-like pocket which are lined with hydrophobic residues and which comprise a zinc binding site, several polar side chains, including two Asp-His charge

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relay systems. Mutation of the zinc ligands and other polar residues at the pocket bottom reduces or eliminates the catalytic activity.

The present inventors found that mutation at the Tyr297Phe site reduced activity. See also, Hassig et al., 1998, Proc. Natl. Acad. Sci. USA 95:3519-3524; Kadosh & Struhl, 1998, Genes Dev. 12:797-805. The elimination of activity by mutation of these residues indicates that this region is the enzyme active site. Adjacent to the active site, there is an internal cavity that may provide space for the diffusion of the acetate reaction product. Homology at the active site between HDLP and HDAC1, as described above, indicates that they share structural and functional homology.

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The inhibitor compound, trichostatin A (TSA) (Tsuji et al., 1976, J. Antibiotics 29:1-6) binds HDLP by inserting its long aliphatic chain, which has a hydroxamic acid group at one end, into the pocket (Figure 6A, 6B and 6C). The aliphatic chain makes multiple contacts in the well-like, hydrophobic portion of the pocket. The hydroxamic acid reaches the polar bottom of the pocket, where it coordinates the zinc in a bidentate fashion and also forms hydrogen bonds with the polar residues in the active site, including the two charge relay system histidines. The aromatic dimethylamino-phenyl group at the other end of the TSA chain makes contacts at the pocket entrance and serves to cap it. The amino acid residues of HDLP which contact TSA are conserved in HDAC, indicating that TSA binds and inhibits HDAC in a similar fashion to HDLP.

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In the complex, the hydroxamic acid, most of the aliphatic chain and part of the dimethylamino-phenyl group of TSA are buried (60% of TSA's surface area; Figure 6A). The hydroxamic acid group binds the zinc in a bidentite fashion forming bonds through its carbonyl (2.4 Å) and hydroxyl groups (2.2 Å) resulting in a penta-coordinated Zn²⁺ (Figure 6B and 6C). The hydroxamic acid hydroxyl group replaces the water molecule that binds to the zinc in the apo-HDLP structure described above. The hydroxamic acid also hydrogen bonds with both charge relay system histidines (hydroxyl oxygen to His131 Ne2, 2.8 Å; and nitrogen to His132 Ne2, 2.8 Å), and the Tyr297 hydroxyl group (2.4 Å; Figure 6B and 6C).

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15 The 5-carbon long branched alkene chain of TSA fits snugly in the narrow portion of the pocket making multiple van der Waals contacts with all of the hydrophobic groups lining the pocket (Figure 6B and 6C). Near its center, the chain contains a methyl substituted carbon-carbon double bond 20 which is sandwiched between the phenyl groups of the Phe141 and Phe98 at the tightest point of the pocket (Figure 6A and The length of the alkene chain appears optimal for spanning the length of the pocket, and allowing contacts both at the bottom and at the entrance of the pocket, although, the cap group of Formula (I) may provide length to 25 span the pocket allowing for a shorter alkene chain (aliphatic chain).

At the entrance of the pocket, one face of the planar structure formed by the dimethylamino-phenyl and adjacent carbonyl groups of TSA makes contacts at the rim of the pocket (Pro22, Tyr91, Phel41; Figure 6B and 6C). This

packing is facilitated by the roughly 110° angle in the overall structure of TSA at the junction of the aliphatic chain and the dimethylamino-phenyl group (occurring at the sp³ hybridized C8 carbon). Upon TSA binding, the side chain of Tyr91, which is mostly solvent exposed, changes conformation to make space for the dimethylamino-phenyl group. This is the only change near the active site observed upon TSA binding.

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The hydroxamic acid group is a common motif in zinc 10 metalloprotease inhibitors. See U.S. Patent No. 5,919,940 and 5,917,090; See also, Grams et al., 1995, Biochemistry <u>34</u>:14012-14020; Lovejoy et al., 1999, Nat. Struct. Biol. 6:217-221; and Holmes & Matthews, 1981, Biochemistry 20:6912-6920. Like TSA, these inhibitors also coordinate 15 the active site zinc in a bidentate fashion using their hydroxamate hyroxyl and carbonyl oxygens, replace the nucleophilic water molecule with their hydroxamate hydroxyl groups and form hydrogen bonds to the general base (Grams et al., 1995, Biochemistry 34:14012-14020; Lovejoy et al., 20 1999, Nat. Struct. Biol. 6:217-221; and Holmes & Matthews, 1981, Biochemistry 20:6912-6920).

SAHA, which has a ~30-fold weaker inhibitory activity than

TSA (Richon et al., 1998, Proc. Natl. Acad. Sci. USA

95:3003-3007), binds HDLP similarly to TSA (see, e.g.,

Figure 4D). The SAHA hydroxamic acid group makes the same

contacts to the zinc and active site residues, and the

importance of these interactions is underscored by the loss

of activity of SAHA derivatives lacking the hydroxamic group

(Richon et al., 1998, Proc. Natl. Acad. Sci. USA 95:3003-

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3007). The six-carbon long aliphatic chain of SAHA packs in the tube-like hydrophobic portion of the pocket. Compared to TSA however, SAHA's aliphatic chain packs less snugly and makes fewer van der waals contacts, in part, because SAHA lacks TSA's C15 methyl group branch. SAHA also lacks TSA's double bonds in this region, and this may lead to increased flexibility of the aliphatic chain. The cap group of SAHA consists of a phenyl-amino ketone group. In the crystal structure, the phenyl group has weak electron density, suggesting that it does not pack as well as the cap group of TSA. This may be due to the larger separation between the hydroxamic and cap groups of SAHA compared to TSA (compare TSA, Formula (II) and SAHA, Formula (III), below).

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25 (III)

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The determination of the structure of HDLP and HDLP bound to an inhibitory compound has enabled, for the first time, the identification of the active site of HDLP and of related HDLP proteins, such as proteins belonging to the HDAC family.

The three-dimensional structural information and the atomic coordinates associated with said structural information of HDLP bound to an inhibitory compound is useful in rational drug design providing for a method of identifying inhibitory compounds which bind to and inhibit the enzymatic activity of HDLP, HDAC family proteins and other histone deacetylaselike proteins related to HDLP. Said method for identifying said potential inhibitor for an enzyme comprising deacetylase activity comprises the steps of (a) using a three-dimensional structure of HDLP as defined by its atomic coordinates listed in Figure 16 to 19; (b) employing said three-dimensional structure to design or select said inhibitor; (c) synthesizing said potential potential inhibitor; (d) contacting said potential inhibitor with said enzyme in the presence of an acetylated substrate; and (e) determining the ability of said inhibitor to inhibit said deacetylase activity.

The potential HDLP and HDLP-related (e.g. HDAC) inhibitors identified by the method of the present invention are represented by formula (I)

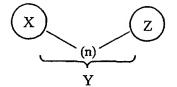
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(I)



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wherein X comprises a cap group which binds to at least one amino acid selected from the group consisting of proline and leucine; Y comprises an aliphatic chain group which binds to at least one amino acid selected from the group consisting of leucine, phenylalanine and glycine; and Z comprises an active site binding group which binds to at least one amino acid selected from the group consisting of aspartic acid, tyrosine and histidine and wherein Z may further bind to a zinc atom and with the provision that the compound of Formula (I) is not TSA, trapoxin, SAHA, SAHA derivatives described in U.S. Patent Nos. 5,608,108; 5,700,811; 5,773,474; 5840,960 and 5,668,179.

The present invention permits the use of molecular design techniques to design, identify and synthesize chemical entities and compounds, including inhibitory compounds, capable of binding to the active site of HDLP and HDLP-related proteins. The atomic coordinates of apo-HDLP and inhibitor-bound HDLP may be used in conjunction with computer modeling using a docking program such as GRAM, DOCK, HOOK or AUTODOCK (Dunbrack et al., 1997, Folding & Design 2:27-42) to identify potential inhibitors of HDLP and HDLP-related proteins (e.g. HDAC1). This procedure can include computer fitting of potential inhibitors to the active site of HDLP to ascertain how well the shape and the

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chemical structure of the potential inhibitor will complement the active site or to compare the potential inhibitors with the binding of TSA or SAHA in the active See Bugg et al, 1998, Scientific American December:92-98; West et al., 1995, TIPS <u>16</u>:67-74. The potential inhibitors designed by modeling with a docking program conform to the general formula (I) as described Computer programs may also be employed to estimate above. the attraction, repulsion and stearic hindrance of the HDLP and potential inhibitor compound. Generally, the tighter the fit, the lower the stearic hindrances, the greater the attractive forces, and the greater the specificity which are important features for a specific inhibitory compound which is more likely to interact with HDLP and HDLP-related proteins rather than other classes of proteins. features are desired particularly where the inhibitory compound is a potential antitumor drug.

The compounds of the present invention may also be designed by visually inspecting the three-dimensional structure to 20 determine more effective deacetylase inhibitors. This type of modeling may be referred to as "manual" drug design. Manual drug design may employ visual inspection and analysis using a graphics visualization program such as "O" (Jones, T.A., Zhou, J.Y., Cowan, S.W., and Kjeldgaard, M., Improved 25 method for building protein models in electron density maps location of errors and the in these models, Acta Crystallog., A47, 110-119.

Initially potential inhibitor compounds can be selected for their structural similarity to the X, Y and Z constituents

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- 41 -

of formula (I) by manual drug design. The structural analog thus designed can then be modified by computer modeling programs to better define the most likely effective candidates. Reduction of the number of potential candidates is useful as it may not be possible to synthesize and screen a countless number of variations compounds that may have some similarity to known inhibitory molecules. analysis has been shown effective in the development of HIV protease inhibitors (Lam et al., 1994, Science 263:380-384; Wlodawer et al., 1993, Ann. Rev. Biochem. 62:543-585; Appelt, 1993 Perspectives in Drug Discovery and Design 1:23-48; Erickson, 1993, Perspectives in Drug Discovery and Design 1:109-128. Alternatively, random screening of an small molecule library could lead to potential inhibitors whose inhibitory activity may then be analyzed by computer modeling as described above to better determine their effectiveness as inhibitors.

The compounds designed using the information of the present invention may be competitive or noncompetitive inhibitors. These designed inhibitors may bind to all or a portion of the active site of HDLP and may be more potent, more specific, less toxic and more effective than known inhibitors for \mathtt{HDLP} and HDLP-related proteins, particularly HDACs. The designed inhibitors may also be less potent but have a longer half life in vivo and/or in vitro and therefore be more effective at inhibiting histone deacetylase activity in vivo and/or in vivo for prolonged periods of time. Said designed inhibitors are useful to inhibit the histone deacetylase activity of HDLP and HDLPrelated proteins (e.g. HDAC1), to inhibit cell growth in

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vitro and in vivo and may be particularly useful as antitumor agents.

The present invention also permits the use of molecular design techniques to computationally screen small molecule data bases for chemical entities or compounds that can bind to HDLP in a manner analogous to the TSA and SAHA as defined by the structure of the present invention. computational screening may identify various groups which may be defined as "X", "Y" or "Z" of formula (I) above and may be employed to synthesize the potential inhibitors of the present invention comprising formula (I). potential inhibitors may be assayed for histone deacetylase inhibitory activity in a histone deacetylase activity assay (see Example 3 below), may be co-crystallized with HDLP to determine the binding characteristics through X-ray crystallography techniques defined above (e.g. said cocrystal structure may be determined by molecular replacement to assess the binding characteristics of said potential inhibitor), or may be assessed based on binding activity by incubating said potential inhibitor with said HDLP, performing gel filtration to separate any free potential inhibitor to HDLP-bound inhibitor, and determining the amount of histone deacetylase activity of the inhibitorbound HDLP. To measure binding constants (e.g., Kd), methods known to those in the art may be employed such as $\operatorname{Biacore}^{\operatorname{m}}$ analysis, isothermal titration calorimetry, Elisa with a known drug on the plate to show competitive binding, or by a deacetylase activity assay.

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The design of potential inhibitors of the present invention is further facilitated by reference to Figure 9, which is a surface representation figure that depicts the surface grooves. Analysis of such grooves gives insight into the constituents of the cap group of formula (I). The surface grooves are labeled groove A, groove A', groove B and groove C, into which additional cap groups may bind. The structure of HDLP bound to either TSA or SAHA shows that the cap groups of TSA and SAHA bind in groove A. By analysis of the amino acid sequence identity of HDLP and HDACs, Groove A is well conserved in HDACs, has a significant hydrophobic component, appears deep enough to allow for significant interactions and is also the largest of the four grooves. In addition to the dimethylamino phenyl group of the TSA, the A groove can fit approximately 200 daltons worth of groups (e.g. groove A could accommodate a naphthalene-like group after an appropriate spacer, etc.). Groove A. as referred to herein, is characterized by the following conserved residues of HDLP: His 21, Pro22, Lys24, Phe141, Leu265 and Phe335. The periphery of groove A comprises unconserved residues. Additionally, Groove A', as referred to herein, comprises primarily unconserved residues.

Groove B is immediately adjacent to the pocket. Of significance is that the bottom of groove B comprises the N-epsilon nitrogen of His170, which coordinates the zinc through its N-delta nitrogen. Significant binding energy may be achieved by contacting the Ne proton of His170 with a carboxylic acid or sulfate group. In addition, groove B may be large enough to fit a phenyl group, the face of which may comprise a partial negative charge which may pack over the N-epsilon proton of His170. The conserved residues of

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groove B, as referred to herein are: His170, Tyr196 and Leu265.

Groove C is not as well conserved as the other two grooves and the amino acid residues which comprise groove C are mostly polar and solvent exposed. Groove C, as referred to herein comprises the following conserved residues: Asn87, Gly140 and Phe198.

The compounds of the present invention are represented by formula (I):

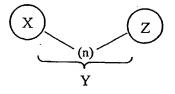
(I)

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Examples for suitable X constituents wherein X comprises a cap group may be described in three categories, depending upon which surface of groove A, A', B and/or C they are targeted to. The cap group may comprise all three categories on the same compound. Of particular benefit may be replacing the cap group of TSA or SAHA with a large, rigid structure. Nonlimiting examples for suitable cap groups (X) of formula (I) which may bind in groove A are: (1) attaching a 1-3 methyl linker followed by a phenyl or naphthalene group from the para or meta position of SAHA's phenyl group represented by formula (IV):

- 45 **-**

(IV)

$$n(H_2C)$$

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(2) attaching a 2-3 methyl linker followed by a phenyl or naphthalene group from the meta position of TSA's phenyl cap group, or from TSA's dimethyl amino group represented by formula (V):

(V)

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and which may bind in groove B is a 1-3 methyl group spacer followed by a carboxylate, sulfate or phenyl group as represented by formula (VI):

(VI)

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With respect to the aliphatic (Y) group, the diameter of the pocket suggests that one more methyl "side chain" could fit, in addition to the C15 methyl group on the C10 carbon. Nonlimiting suitable examples for Y constituents wherein Y comprises an aliphatic chain group are as follows: (1) add

a methyl group to TSA on the C12 carbon (with or without a methyl group on the C10 carbon and with or without double bonds and with or without substituting the X and/or Zconstituents of formula (I)as represented by formula (VII):

(VII)

(2) add a methyl group to TSA on the C9 carbon (with or 10 without a methyl group on the ClO carbon; with or without both or either of the double bonds, and with or without substituting the X and/or Z constituents of formula (I) as represented by formula (VIII):

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(VIII)

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(3) replace the two alkalene double bonds of TSA with only one between C10 and C11, which may free the C11 and C12 torsion to allow for a better fit, the X and/or Z groups may also be substituted as represented by formula (IX):

25 (IX)

(4) cyclize C15 and C12 carbons of TSA through a sulphur atom (or nitrogen atom), the X and/or Z groups may also be substituted as represented by formula (X):

(X)

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(5) extend from the C9 carbon of TSA such that the extension approaches and/or enters groove B (see Figure 9); making C9 sp3 so that it can have some freedom; attach to C9 a 1-3 methyl group spacer which may include a double bond and they attaching thereto a sulfate, carboxylate, sulfate, hyroxyl, or phenyl group which may make an interaction with the N-epsilon proton of Hisl70 which may coordinate the zinc atom as represented by formula (XI):

(XI)

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$$X \xrightarrow{14} \begin{cases} 15 \\ 10 \\ 11 \end{cases}$$

$$Z \xrightarrow{12} R1 = \begin{cases} -COOH \\ -SO_4 \\ -OH \end{cases}$$

$$R_1 = \begin{cases} -COOH \\ -SO_4 \\ -OH \end{cases}$$

25 (6) extend off the C8 carbon (replacing C14) of TSA such that the extension approaches or enters groove B; attach a 1-3 methyl group spacer (which may include a double bond) and then link thereto a carboxylate, sulfate, hydroxyl or phenyl group such that an interaction is made with the N-epsilon proton of His170 that coordinates the zinc atom; the X and/or Z constituents may also be substituted as represented by formula (XII):

(XII)

$$\begin{array}{c|c} R_1 \\ CH_2)n & (15) \\ \hline X & R_1 = \begin{cases} -COOH \\ -SO_4 \\ -OH \end{cases} \end{array}$$

(7) substitute the C8 carbon at the end of the aliphatic chain such that the substitution may contact groove A, A', B and or C, in such an example, a cap group (X) may or may not be required and the X and Z constituents may be substituted as well, as represented by formula (XIII):

(XIII)

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(8) formulas VII through XIII above wherein the aliphatic chain further comprises a methyl group between the active site binding group (Z) and the C8 carbon, and preferably just before the C8 carbon, increasing the distance between X and Z, (9) make the connection between the aliphatic chain and the cap group more rigid (e.g., by closing a 6-membered ring which may or may not comprise oxygen, the X and Z group may also be substituted as represented by formula (XIV):

(XIV)

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and (10) combining two or more of the changes depicted by formulas (VII-XIV).

Additionally, nonlimiting examples for suitable Z groups wherein Z comprises an active site binding group are as follows: (1) hydroxamic acid, (2) carboxylic acid, (3) sulfonamide, (4) acetamide, (5) epoxyketone, (6) an ester with a methyl linker and a hydroxyl of acetate ester group to lead into the cavity and interact with a conserved arginine (Arg27) as represented by formula (XV): (XV)

$$R_1 = \begin{cases} CH_2 | h & 0 \\ CH_2 | h & CH_2 | h \end{cases}$$

$$R_1 = \begin{cases} -OH & 0 \\ -CH_2 | h & CH_2 | h \end{cases}$$

$$O - - CH_3$$

and (7) an alphaketone as represented by formula (XVI):
(XVI)

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Additionally, other suitable X, Y and Z constituents may be envisioned by the skilled artisan given the three-dimensional structural information of the present invention.

After having determined potential suitable X, Y and Z constituents, the constituents are combined to form a compound of formula (I) using combinatorial chemistry techniques. This may be achieved according to U.S. Patent Nos. 5,608,108; 5,700,811; 5,773,474; 5,840,960 and 5,668,179, incorporated herein by reference. Any methods

known to one of skill in the art may be employed to synthesize compounds of formula (I) comprising X, Y and Z constituents as determined by the methods described above.

- As mentioned above, the compounds of formula (I) are useful to inhibit the histone deacetylase activity of HDLP and HDAC-related proteins. Such inhibition may allow for a reduction or cessation of cell growth in vitro and in vivo.
- For in vitro use, such reduction or cessation of cell growth 10 is useful to study the role of histone deacetylation and differentiation during the cell cycle and also to study other mechanisms associated with cell cycle arrest and particularly how the repression of transcription is involved in cell cycle progression which may be studies in a yeast 15 model system such as that described by Kadosh & Struhl, 1998, Mol. Cell. Biol. <u>18</u>:5121-5127. In vitro model systems which may be employed to study the effects of potential inhibitors on cell cycle progression and also tumor growth include those described by: Richon et al, 1998, Proc. Natl. 20 Acad. Sci. USA 95:3003-3007; Yoshida et al., 1995, Bioessays 17:423-430; Kim et al., 1999, Oncogene 18:2461-2470; Richon et al., 1996, Proc. Natl. Acad. Sci. USA 93:5705-5708; and Yoshida et al., 1987, Cancer Res. 47:3688-3691.

For in vivo use, such a reduction or cessation of cell growth is useful to study the effect of said inhibitor compounds in non-human animal model systems of cancer and is also useful for the treatment of cancer in a recipient in need of such treatment. Non-limiting examples of animals which may serve as non-human animal model systems include

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mice, rats, rabbits, chickens, sheep, goats, cows, pigs, and non-human primates. See, e.g., Desai et al., 1999, Proc. AACR 40: abstract #2396; Cohen et al., 1999, Cancer Res., submitted. The compounds of the present invention may be administered to a transgenic non-human animal wherein said animal has developed cancer such as those animal models in which the animal has a propensity for developing cancer (e.g. animal model systems described in U.S. Patents 5,777,193, 5,811,634, 5,709,844, 5,698,764, and 5,550,316). Such animal model systems may allow for the determination of toxicity and tumor reduction effectiveness of the compounds of the present invention.

A preferred compound of the present invention may comprise high specific activity for HDLP and HDAC-related proteins, good bioavailability when administered orally, activity in reducing or ceasing cell growth in tumor cell lines, and activity in reducing or ceasing tumor growth in animal models of various cancers.

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Accordingly, another aspect of this invention is a method of eradicating or managing cancer in a recipient, which may be an animal and is preferably a human. Said method comprises administering to said recipient a tumor reducing amount of a compound as defined by formula (I) above, or a physiological acceptable salt thereof.

In a further aspect of the invention, there is provided a composition comprising the compound of formula (I) and an excipient or carrier. Administration of the foregoing agents may be local or systemic. Such carriers include any

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suitable physiological solutions or dispersant or the like. The physiologic solutions include any acceptable solution or dispersion media, such as saline, or buffered saline. The carrier may also include antibacterial and antifungal agents, isotonic and absorption delaying agents, and the like. Except insofar as any conventional media, carrier or agent is incompatible with the active ingredient, its use in the compositions is contemplated.

- Routes of administration for the compositions containing the 10 delivery vehicle constructs of the present invention include any conventional and physiologically acceptable routes, such as, for example, oral, pulmonary, parenteral (intramuscular, intraperitoneal, intravenous (IV) subcutaneous or. injection), inhalation (via a fine powder formulation or a 15 fine mist), transdermal, nasal, vaginal, rectal, sublingual routes of administration and can be formulated in dosage forms appropriate for each route of administration.
- The following examples are provided to more clearly illustrate the aspects of the invention and are not intended to limit the scope of the invention.

EXAMPLES

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25 Example 1: Protein Production and Purification:

Full-length wild-type HDLP (Genbank accession number AE000719) was subcloned from an Aquifex aeolicus chromosomal DNA preparation (provided by Robert Huber of Universitaet of Regensburg, Germany) into the pGEX4T3 (Amersham-Pharmacia, Piscataway, NJ) vector using the polymerase chain reaction (PCR). The cysteine-to-serine and active site mutants were constructed by PCR site directed mutagenesis and were

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sequenced. The HDLP-glutathione S-transferase (GST) fusion protein was produced in Escherichia coli, purified by affinity chromatography using a column glutathione-sepharose resin (Amersham-Pharmacia, Piscataway, NJ), and by anion-exchange chromatography (Q-sepharose™; Amersham-Pharmacia, Piscataway, NJ). HDLP was cleaved from the fusion protein with thrombin at 4°C, was purified by anion-exchange (Q-sepharose™; Amersham-Pharmacia, Piscataway, (UN and gel filtration chromatography (Superdex 200; Amersham-Pharmacia, Piscataway, NJ), and was concentrated to typically 25 mg/ml in a buffer of 25 mM bis-tris propane (BTP), 500 mM NaCl, 5 mM dithiothrietiol (DTT), 2% isopropanol, pH 7.0.

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Although, it is not known what metal cofactor HDLP contains 15 in vivo, it is presumed to be zinc because of arrangement of the ligands and the similarities in the active site to the zinc proteases. The lack of metal in the purified HDLP is presumed due, in part, to the use of DTT 20 during purification. HDLP was reconstituted with Zn2+ by mixing the Cys75Ser/Cys77Ser double mutant at 10 mg/ml with a 5-fold molar excess of ZnCl₂ in a buffer of 25 mM bis-tris propane, 200 mM NaCl, 1% isopropanol, pH 7.0. Unbound ZnCl² was removed by fractionating HDLP through a G25 desalting 25 column (Amersham-Pharmacia, Piscataway, NJ). The $\mbox{HDLP-Zn}^{2+}\mbox{-TSA}$ complex was prepared by incubating the \mbox{Zn}^{2+} reconstituted HDLP mutant with 1 mM TSA for 45 minutes, followed by gel filtration chromatography (Superdex™200; Amersham-Pharmacia, Piscataway, NJ) to remove excess TSA, and concentration to typically 25 mg/ml in a buffer of 25 mM 30 bis-tris propane, 500 mM NaCl, 1% isopropanol, pH 7.0.

FLAG epitope tagged human HDAC1 was overexpressed using a baculovirus expression system in Hi5 (Invitrogen, Carlsbad, CA) insect cells grown in suspension in serum-free media (Sf900, Gibco, Grand Island, NY). The fusion protein was purified by anion exchange and affinity chromatography using Anti-FLAG M2 affinity resin (Sigma, St. Louis, MO) and FLAG Peptide (Sigma,, St. Louis, MO).

Example 2: Crystallization and data collection:

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10 Crystals of apo-HDLP were grown at room temperature by the hanging-drop vapor-diffusion method, from 7.5% isopropanol, 28% PEG 1500, 425 mM NaCl, 100 mM Tris-Cl, pH 7.0. They form in space group C2 with a = 51.4 Å, b = 93.8 Å, c = 78.7 Å, β = 96.9 Å, and contain one HDLP molecule in the asymmetric unit. Diffraction data were collected with crystals flash-frozen in a buffer of 7.5% isopropanol, 35% PEG 1500, 75 mM NaCl, 100 mM Tris-Cl, pH 8.0, at -170° C.

The structure of the HDLP- Zn²⁺ complex was determined from HDLP Cys75Ser/Cys77Ser double mutant crystals grown from 23% tert-butanol, 27% PEG 1500, 400 mM KCl, 100 mM bis-tris propane-Cl, pH 6.8. Space group and cell dimensions were identical to the apocrystals. The HDLP-Zn²⁺ crystals were harvested and frozen in 27% tert-butanol, 22% PEG 1500, 50 mM KCl, 20 mM NaCl, 0.2 mM ZnCl₂, 100 mM bis-tris propane, pH 6.8, at -170° C.

Crystals of the HDLP-Zn²⁺-TSA complex comprised HDLP Cys75Ser/Cys77Ser double mutant and were grown from 23% tert-butanol, 27% PEG 1500, 600 mM KCl, 100 mM bis-tris propane-Cl, pH 6.8, by microseeding. The crystals were grown in the presence of zinc. They form in space group

 $P2_12_12_1$ with a = 53.4 Å, b = 94.4 Å, c = 156.3 Å and contain two HDLP- Zn^{2+} -TSA complexes in the asymmetric unit. The HDLP- Zn^{2+} -TSA crystals were harvested and frozen in the same cryobuffer as the HDLP- Zn^{2+} crystals except that 0.5mM TSA was added. Data were processed with DENZO and SCALEPACK (Otwinowski & Minor, 1997, Method. Ensemble. 276:307-326). MIR analysis, model building and refinement.

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The HDLP-Zn²⁺-SAHA complex crystals were grown and evaluated the same as the HDLP-Zn²⁺-TSA crystals. However, the restraints for the SAHA structure were constructed based on stereochemical parameters from TSA. Like the apo-HDLP crystals, the SAHA/HDLP co-crystals grew in space group C2.

15 Heavy-atom soaks were performed with the apo-HDLP crystals in a buffer of 7.5% isopropanol, 30% PEG 1500, 75 mM NaCl, 100 mM Tris-Cl, pH 8.0, supplemented with 1.0 mM thimerosal for 2h, 5 mM KAu(CN)₂ for 1h, and 1 mM Pb(Me)₃OAc for 2h. MIR phases were calculated with the program MLPHARE (The CCP4 suite: Programs for computational crystallography, 20 1994, Acta Crystallogr. D 50:760-763) at 2.5 Å using the anomalous diffraction signal from the thimerosal derivative, and had a mean figure of merit of 0.55. The phases were improved by solvent flattening with the program DM (The CCP4 suite: Programs for computational crystallography, 1994, 25 Acta Crystallogr. D 50:760-763) , and were used to build the initial model with the program O (Jones et al., 1991, Acta Crystallogr. A 47:110-109). Successive rounds of rebuilding and simulated annealing refinement with the program CNS (Brunger et al., 1998, Acta Crystallogr. D 30 54:905-921) allowed interpretation of HDLP from residues 2

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to 373. Residues 1, 374, and 375 were not modeled and are presumed to be disordered.

The structure of the HDLP- $\mathrm{Zn^{2+}}$ -TSA and HDLP- $\mathrm{Zn^{2+}}$ -SAHA complex were determined by molecular replacement with the program AMORE (The suite: Programs for computational CCP4 crystallography, 1994, Acta Crystallogr. D 50:760-763) using the apo-HDLP structure as a search model. The initial electron density maps had strong and continuous difference density for the entire TSA molecule. However the SAHA molecule was not as well ordered in the cap group region. The structure of TSA was obtained from the Cambridge Structural Database (Refcode TRCHST) and was used to define stereochemical restraints used in the refinement with the program CNS. The restraints of SAHA were constructed based on stereochemical parameters from TSA and surrounding amino acid residues. The dimer interface in the $HDLP-Zn^{2+}-TSA$ and $\mbox{HDLP-Zn}^{2+}\mbox{-SAHA}$ crystals primarily involves Phe200 on the protein surface. The Phe200 side chain contacts Tyr91, whose side chain conformation changes on TSA binding, and part of the dimethyl amino phenyl group of TSA from the second protomer. The HDAC family does not contain a phenylalanine residue at the equivalent position.

25 Example 3: Histone deacetylase assays:

Purified proteins were assayed by incubating 10 μ g of [³H]acetyl-labeled murine erythroleukemia histone substrate and HDAC assay buffer (20 mM Tris-HCl, pH 8.0, 150 mM NaCl, 10% glycerol) for 30-60 minutes at 37° C in a total volume of 30 μ l. The final concentrations of HDLP and HDACl-FLAG were 3.6 μ M and 0.24 μ M, respectively. Assays were performed in duplicate. The reactions were stopped and the

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released acetate was extracted and assayed as described (Hendzel et al., 1991, J. Biol. Chem. 266:21936-21942). [3H] acetyl-labeled murine erythroleukemia histones were prepared essentially as described (Carmen et al., 1996, J. Biol. Chem. 271:15837-15844). Inhibitors were added in the absence of substrate and incubated on ice for 20 minutes, substrate was added, and the assay performed as described above. HDLP was inclubated with 20 μ M ZnCl₂ and 20 μ M MnCl₂(H2O)₄ in HDAC buffer and tested for activity.

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Only HDLP dialyzed against ${\rm ZnCl_2}$ had activity. HDAC1-FLAG was dialyzed against 20 μM ${\rm ZnCl_2}$ in HDAC buffer which had no effect on activity. Therefore, HDAC1-FLAG contains a metal as purified.

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The in vivo substrate of HDLP is not known. HDLP may have a role in acetoin utilization like the B. subtilis AcuC gene product, and it has been annotated as such in the genome sequence, but the reaction catalyzed by AcuC is also not known. Furthermore, the A. aeolicus genome appears to lack the acuA and acuB genes that are part of the acuABC operon of B. subtilis (Deckert et al., 1998 Nature 392:353-358), and HDLP is as similar to human HDAC1 (35.2 % identity) as it is to B. subtilis AcuC (34.7 % identity).

What is claimed is:

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- 1. A crystal of an enzyme comprising deacetylase activity wherein said crystal effectively diffracts X-rays for the determination of the atomic coordinates of said enzyme to a resolution of greater than 4 Å and wherein the structure of said enzyme comprises a conserved core α/β structure characteristic fold wherein said conserved α/β fold comprises an eight-stranded parallel β sheet and eight α helices and wherein four of the helices pack on either face of said parallel β sheet and wherein said structure of said enzyme comprises an rmsd of less than or equal to 1.5 Å in the positions of C α atoms for at least 2/3 or more of the amino acids of HDLP as defined by the atomic coordinates of HDLP.
 - 2. The crystal of claim 1, wherein said protein structure further comprises:
 - (a) eight α helices positioned near one side of the β sheet; and
 - (b) at least seven large, well defined loops originating from the C-terminal ends of the β -strands of said eight-stranded parallel β sheet wherein the eight extra helices and the seven large loops are associated with a significant extension of the structure beyond the core α/β motif and wherein said extension of the structure gives rise to a deep, narrow pocket and an internal cavity adjacent to the pocket.

3. The crystal of claim 1, wherein said enzyme comprising deacetylase activity is selected from the group

consisting of HDLP, HDLP-related proteins, HDAC1, HDAC2, HDAC3, HDAC4, HDAC5, HDAC6, HDAC-related proteins, APAH, AcuC, and functional derivatives thereof.

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- 4. The crystal of claim 2 further comprising a specifically bound zinc atom in the active site of said enzyme.
- 10 5. The crystal of claim 2 further comprising a specifically bound deacetylase inhibitor compound in the active site of said enzyme.
- 6. The crystal of claim 2 define by the atomic coordinates according to Figure 16.
 - 7. A method for identifying a potential deacetylase inhibitor compound for an enzyme which comprises deacetylase activity, said method comprising the steps of:
 - a. using a three-dimensional structure of HDLP as defined by atomic coordinates according to Figure 16;
 - b. employing said three-dimensional structure to design or select said potential inhibitor;
 - c. synthesizing said potential inhibitor;
 - d. contacting said potential inhibitor with said enzyme in the presence of an acetylated substrate; and
- determining the deacetylase inhibitory activity of said potential inhibitor.

- 60 -

- 8. The method of claim 7, wherein the three-dimensional structure is designed or selected using computer modeling.
- 5 9. The method of claim 7, wherein the potential deacetylase inhibitor is designed de novo.
- 10. The method of claim 7, wherein the potential deacetylase inhibitor is designed based on a known inhibitor.
- 11. The method of claim 7, wherein said enzyme comprising deacetylase activity is selected from the group consisting of HDLP, HDLP-related proteins, HDAC1, HDAC2, HDAC3, HDAC4, HDAC5, HDAC6, HDAC-related proteins, APAH, and AcuC.
- 12. A method of evaluating the binding properties of the potential deacetylase inhibitor compound comprising the steps of:
 - a. co-crystallizing said compound with HDLP;

25

- b. determining the three-dimensional structure of said HDLP-potential inhibitor complex co-crystal by molecular replacement using the threedimensional structure of HDLP as defined by atomic coordinates according to Figure 16; and
- c. analyzing said three-dimensional structure of said HDLP bound to said potential inhibitor compound to evaluate the binding characteristics of said potential inhibitor compound.
- 13. A method for solving the structure of an HDAC family

member crystal comprising the steps of:

- a. collecting X-ray diffraction data of said crystal wherein said data diffracts to a high resolution limit of greater than 4 Å;
- b. using the atomic coordinates of HDLP accoding to Figure 16 to perform molecular replacement or refinement and difference fourier with said X-ray diffraction data of said HDAC family member crystal to determine the structure of said HDAC family member; and
 - c. refining said structure of said HDAC family member.
- 14. The method of claim 13, wherein said HDAC family member is HDAC1.
 - 15. A Cys75Ser/Cys77Ser double mutant of HDLP wherein said mutant is encoded by the nucleic acid sequence of SEQ ID NO:4.
 - 16. A Cys75Ser/Cys77Ser double mutant of HDLP wherein said mutant has the amino acid sequence of SEQ ID NO:3.
 - 17. A nucleotide sequence according to SEQ ID NO:4

20

- 18. An expression vector comprising the nucleotide sequence of claim 17.
- 19. A method of using the crystal of claim 1 for screening30 for a novel drug comprising:
 - a. selecting a potential ligand by performing

5.

rational drug design with the three-dimensional structure determined for the crystal;

- contacting the potential ligand with the ligand binding domain of the crystal; and
- c. detecting the binding potential of the potential ligand for the ligand binding domain, wherein the novel drug is selected based on its having a greater affinity for the ligand binding domain than that of a known drug.

Figure 1

Statistics from the crystallographic analysis

TABLE 1.

1,04 3.83

1.48 1.78

0.009

25.8 25.8

22.0. 22.4

434 456

3424 6475

44,122

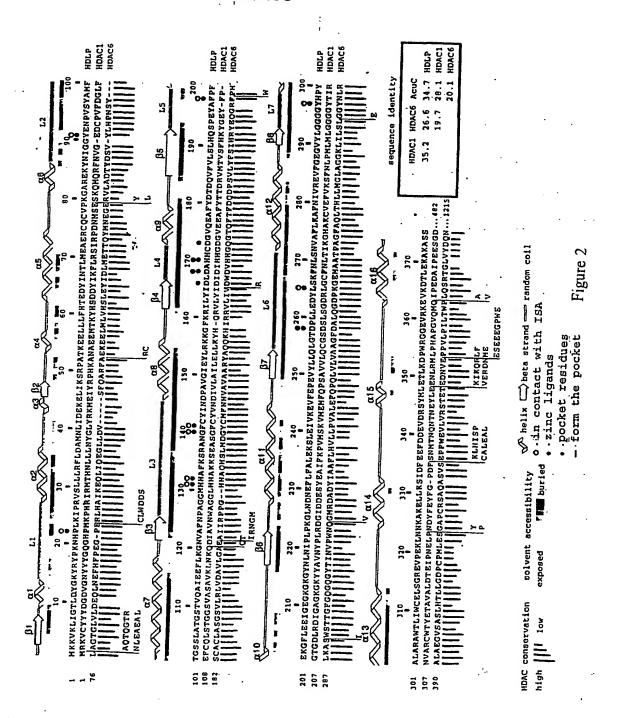
HDLP-Zn-TSA

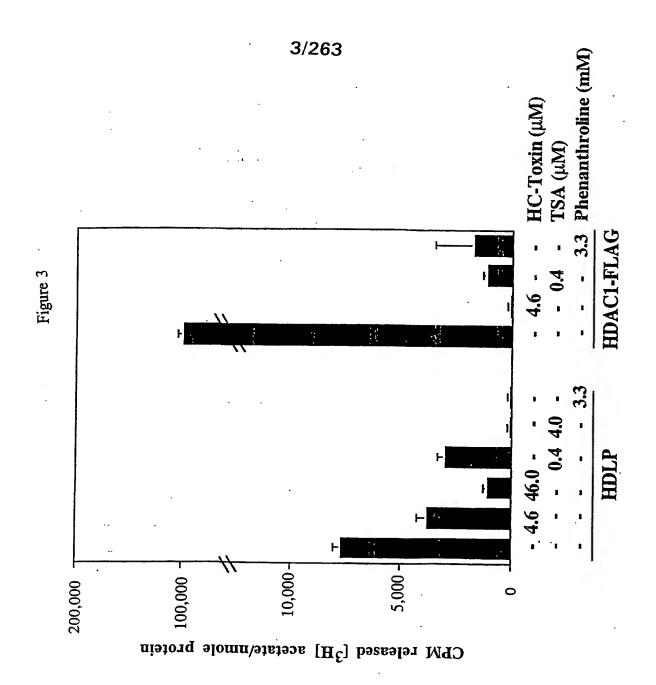
1/263

Data set		Na	Native th	thimerosal	Pb	Aucn	Zu	TSA		
Space Group Resolution (Å) Observations Unique reflections Data coverage (%) Rsym (%) MIR analysis (20.0- phasing power Rcullis Rcullis (ano)	oup (A) nns lections age (%) sis (20.0-2.5 Å); ower		C2 1.8 134,952 32,143 92.3 2.9	C2 2.3 79,023 15,958 95.7 8.4 1.47 0.72	C2 3.5 11,454 4,040 86.4 9.6 1.24 0.78	C2 2.8 27,722 8,753 94.3 8.9 1.10	C2 2.0 125,769 23,643 90.6 7.2	P2,2,2, 2.1 180,427 50,796 93.8 7.1	3 H	
Refinement statistics:				·			•	RMSD		
Data Set HDLP HDLP-Zn	Resolution F (Å) 1.8 2.0	Reflections (IFI > 1σ) 31,550 23,582	Total atoms 3214 3424	Water atoms 228 434	R-factor (%) 19.8	R-free (%) 24.0	bonds (Å) 0.010	angles (°) 1.63	B-factor (Ų) 3.55	

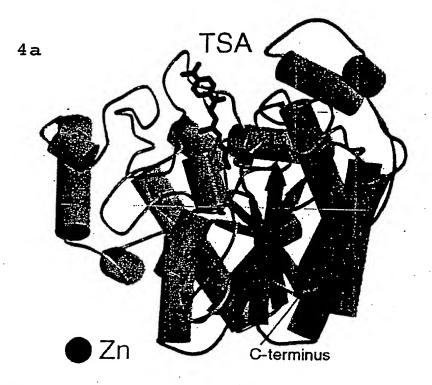
omitted from the start of refinement. RMSD: root mean square deviations from ideal geometry and root where <F $_{\lambda l}>$ is the root-mean-square heavy atom structure factor and E is the residual lack of closure error. Rcullis is the mean residual lack of closure error divided by the dispersive difference. R-factor = $\Sigma | F_{obs}$ - $\mathsf{F}_{\mathsf{calc}}[l\Sigma|\mathsf{F}_{\mathsf{obs}}]$, where $\mathsf{F}_{\mathsf{obs}}$ and $\mathsf{F}_{\mathsf{calc}}$ are the observed and calculated structure factors, respectively. Figure of merit = IF(hkl)best[/F(hkl). R-free = R-factor calculated using 5% of the reflection data chosen randomly and $Rsym = \Sigma_h \Sigma_i \ln_i - < \ln_i > \sum_h \Sigma_i \ln_i$ for the intensity (I) of i observations of reflection h. Phasing power = $< F_{\lambda i} > / E$, mean square variation in the B-factor of bonded atoms.

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4b

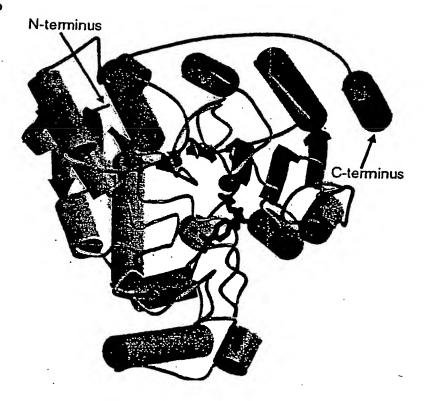
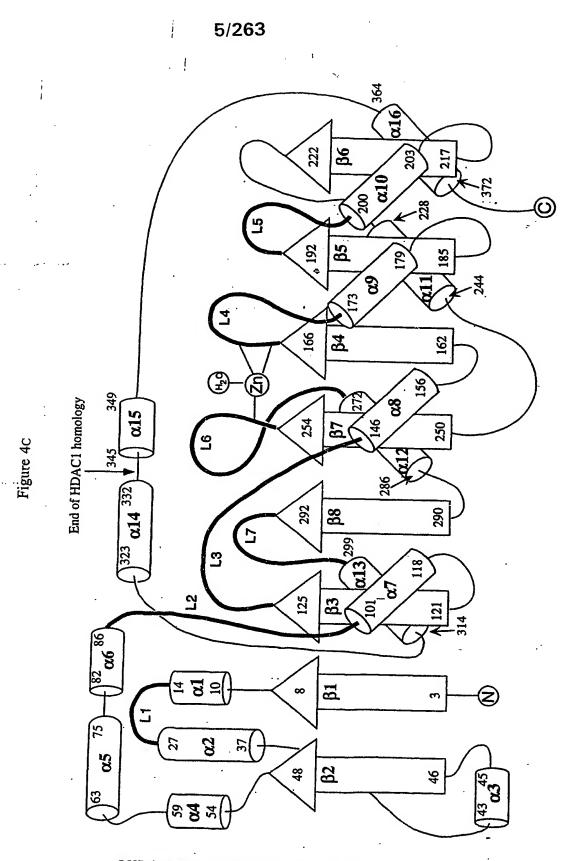


Figure 4
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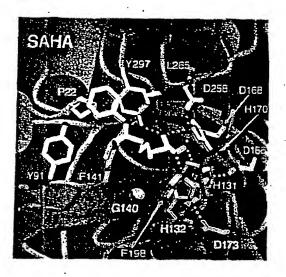
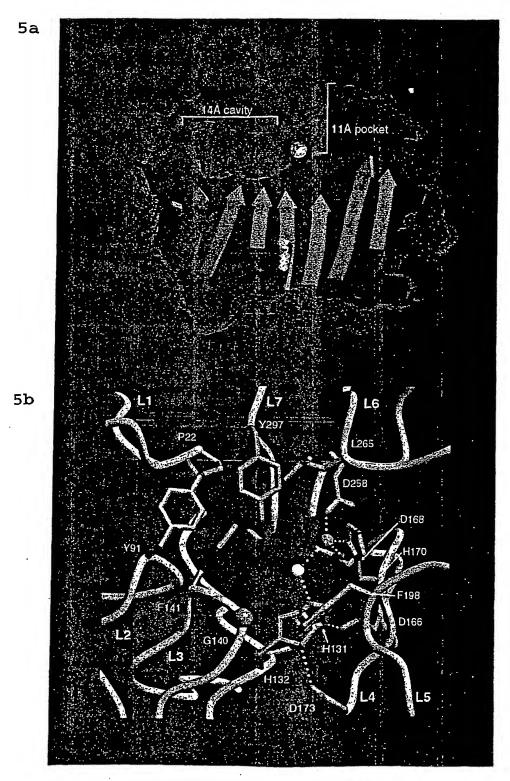


Figure 4D

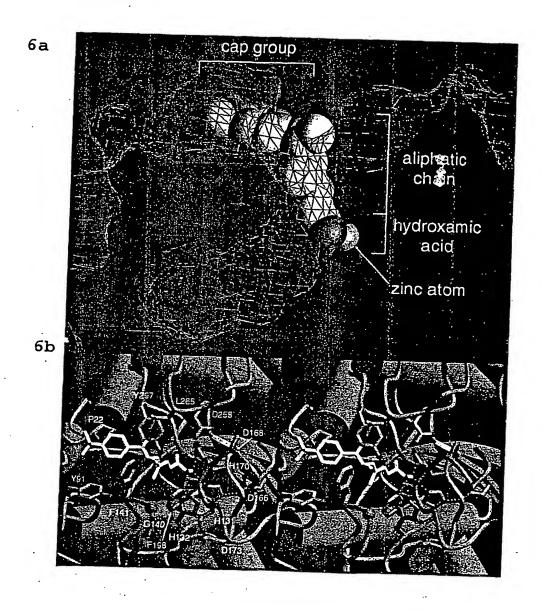
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Figure 6



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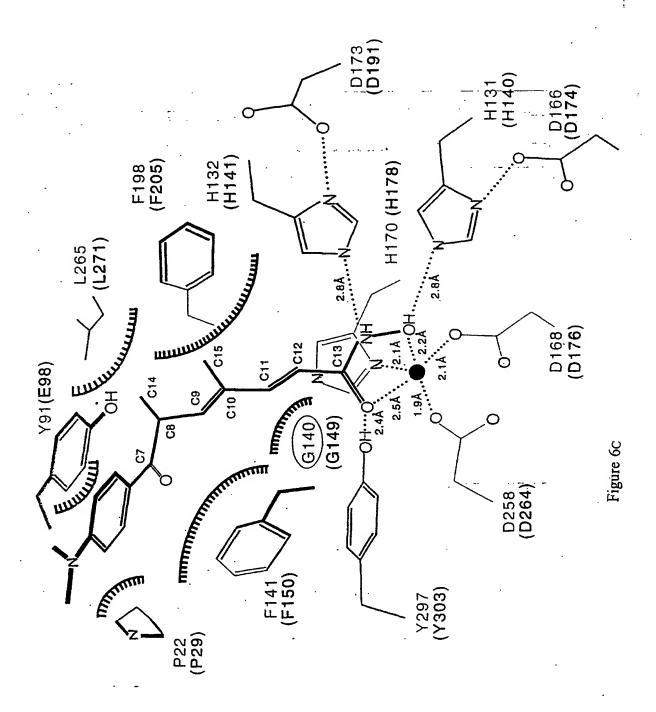
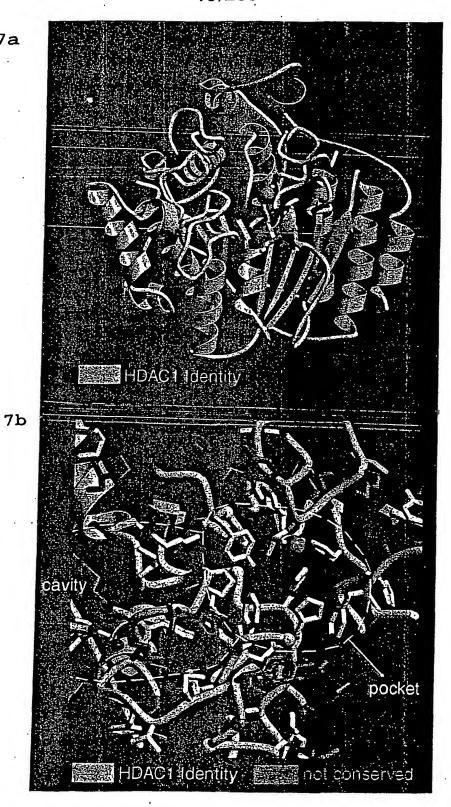


Figure 7

7a



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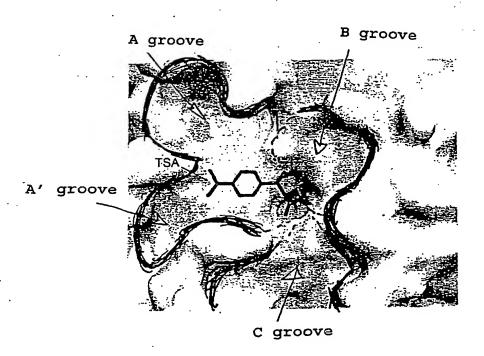


Figure 9

Figure 10

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10 20 30 40 ATGAAGAAGGITAAACTTATCGGAACTTTAGACTACGGAA 40 AGTACAGATATCCCAAAAACCATCCTCTTAAAATACCAAG 80 AGTITICCCTACTCCTTAGGTTTTTAGATGCCATGAACCTT 120 ATAGATGAGAAGGAATTAATCAAGAGCAGACCCGCAACTA 160 AAGAAGAACTCCTTTTATTCCACACGGAAGACTACATAAA 200 210 220 230 240 CACTITAATGGAAGCGGAAAGGIGICAGIGCGIICCGAAG 240 GGAGCTAGGGAAAAGTACAACATAGGCGGATACGAAAACC 280 CCGTATCTTACGCGATGTTTACAGGCTCTTCTCTCGCAAC 320 GGGITCAACAGIGCAGGCGATAGAGGAAITIITTAAAGGGA 360 AATGTAGCTTTCAATCCCGCGGGAGGTATGCACCACGCTT 400 410 420 430 440 TTAAAAGCAGGCAAACGGCTTTTGCTACATAAACGACCC 440 CGCTGTGGGAATTGAGTACTTGAGAAAAAAAGGCTTTAAG 480 AGAATACTCTACATAGACCTTGATGCCCACCACTGCGACG 520 GIGITCAGGAAGCCITTIACGATACAGACCAGGIGITCGI 560 CCIGICCCITCACCAGICGCCGAGIACGCCITICCCITT 600 610 620 630 640 GAGAAGGCTTCCTGGAGGAGGAGGAAAAGGAA 640 AGGCTACAACCTGAACATTCCCCTGCCAAAGGCCTTGAA 680 CGACAACGAGTTCCTCTTTGCCCTAGAAAATCTCTGGAA 720 ATAGTCAAAGAAGTATTTGAGCCCGAGGTTTACCTTCTTC 760 AACTCGGAACTGACCCACTCCTTGAAGATTACCTTTCCAA 800 810 820 830 840 GITCAACCICTCAAACGITGCCITTITTAAAAGCTTTCAAC 840 ATCGITCGTGAGGTTTTCGGGGAGGGAGTATACCTCGGAG 880 GAGGCGGATACCATCCTTACGCCCTCGCAAGGGCATGGAC 920 CCTAATCTGGTGCGAGCTTTCGGGAAGGGAAGTGCCGGAA 960 AAGCTAAACAATAAAGCAAAAGACCTTTTAAAGAGTATAG 1000 1010 1020 1030 1040 resident de la contraction de ACTTIGAAGAGTTIGACGACGAGGTGGACCGCTCGTACAT 1040 GCTCGAAACCCTAAAGGACCCCTGGAGAGGAGGAGGGAGAGGTA 1080 AGGAAAGAAGTAAAGGATACGCTTGAAAAGGCGAAAGCCT 1120 CATCITA 1127

Figure 11

10	20	30	40	
mandanal	<u> </u>	سيلسن	لسيا	
MKKVKLIGILDYGKY	RYPKNHPLK	IPRVSLLLRF	LDAMNL	40
IDEKELIKSRPATKE				80
GAREKYNIGGYENPV	SYAMFTGSS	LATGSTVQAI	EEFLKG	120
NVAFNPAGGMHHAFK	SRANGFCYI	NDPAVGIEYI	RKKGFK	160
RILYIDLDAHHCDGV	QEAFYDTDQ	VFVLSLHQSE	EYAFPF	200
210	220	230	240)
<u> Lintrolinial</u>	سيليير	سيلسب	لسبل	
EKGFLEEIGEGKGKG	YNLNIPLPK	GLNDNEFLFA	LEKSLE	240
IVKEVFEPEVYLLQL	GIDPLLEDY	LSKFNLSNVA	FLKAFN	280
IVRÈVEGEGVYLGGG				320
KLNNKAKELLKSIDF	EEFDDEVDR	SYMLETLKDE	WRGGEV	360
RKEVKDILEKAKASS	275			

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Figure 12

10	. 20	30	40
		سيليب	
ATGAAGAAGGITAA			
AGTACAGATATCCC			
AGTITICCCTACTCC			
ATAGATGAGAAGGA	ATTAATCAAG	AGCAGACCCGC	CAACTA 160
AAGAAGAACTCCTT	ITATTCCACA(CGGAAGACTAC	CATAAA 200
210	220	230	240
<u> </u>	<u>Limber</u>	لتتناييتنا	لبييا
CACTITAATGGAAG	CGAAAGGIG.	ICAGIGCGI'I	CCGAAG 240
GGAGCTAGGGAAAA	TACAACATA(GCGGATACG?	AAAACC 280
CCGTATCTTACGCG	ATGITTACAG	ETCITCICIO	CCAAC 320
GGGTTCAACAGTGC	AGGCGATAGA(GAATTTTAA	AAGGGA 360
AATGTAGCTTICAA			
410	420	430	440
410			
TTAAAAGCAGGCA			
CGCTGTGGGAATTG			
AGAATACTCTACATA			
GIGITCAGGAAGCC			
CCTGTCCCTTCACC			
		•	
610	620	630	640
			•
GAGAAGGCTTCCTC			
AGGCTACAACCTG			
CGACAACGAGITCC		·	• •
ATAGICAAAGAAGIZ			
AACTCGGAACTGACC	CACICCITG	AAGATTACCIT	TCCAA 800
810	820	830	840
<u> ,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,</u>			
GITCAACCICICAAA	CGTTCCCTTT	TTAAAAGCTT	TCAAC 840
ATCGTTCGTGAGGTT			
GAGGCGGATTCCATC			
CCTAATCTGGTGCGAC			· · ·
AAGCTAAACAATAAA		•	
•			•
1010 	1020	1030	1040
ACTITIGAAGAGTITIGA			
GCTCGAAACCCTAAAC			
AGGAAAGAAGTAAAGC			
CATCTTA 1127	SUBSTITI	ITE SHEET	(DHIF 941
	:- Gast 4	os Villabi	INOPE TO

Figure 13

	10	20	30	40	•
سلسلس		سيلسنا	سلسب	بلىسىل	
MKKVKLI	GILDYGK	YRYPKNHPLK	TPRVSLLLR	FLDAMNL	40
IDEKELI	KSRPATK	EELLFHTE	YINILMEAE	RCQCVPK	30
GAREKYN	IGGYENP	VSYAMFIGSS	LATGSTVQA	IEEFLKG	120
NVAFNPA	GCMHHAFI	KSRANGFCYI	NDPAVGIEY	LRKKGFK	160
RILYIDL	DAHHCDG	VQEAFYDIDÇ	VFVLSLHQS	PEYAFPF	200
	210	. 220	230	240)
	·	.220 LL)
-	سسلت		سلسنا	للسلل	
EKGFLEE	IGEGKGK	سلست	GINDNEFLE	ALEKSLE	
EKGFLEE IVKEVFE	IGEGKGK(PEVYLLQI	GYNLNIPLPK	GLNDNEFLF LSKFNLSNV	ALEKSLE AFLKAFN	240 280
EKGFLEE IVKEVFE IVREVFG	IGEGKGK(PEVYLLQI EGVYLGG	GYNLNIPLPK LGIDPLLEDY	GLNDNEFLF ISKFNLSNV AWILIWCEL	ALEKSLE AFLKAFN SGREVPE	240 280

17/263 Figure 14

	10	20	30	40
سيليب	سلسل	سلسبلب	سلسلت	لم
ATGAAGAAC	GITAAACII	ATCGGAACTI	TAGACTACGG	AA 40
AGTACAGAT	ATCCCAAAA	ACCATOCICI	TAAAATACCA	AG 80
AGITICCCI	ACTCCTTAG	GITTTTACAT	TOCCATGAACC	TT 120
ATAGATGAG	AAGGAATTA	ATCAAGAGCA	GACCCGCAAC	TA 160
AAGAAGAAC	TCCTTTTAT	TCCACACGG	AGACTACATA	AA 200
2	10	220	230	240
سيلين	سلسب	سلسبار	سيليسك	1
CACTITAAT	GGAAGCGGA	AAGGAGTCAC	GAGCGTTCCGA	AG 240
GGAGCTAGG	GAĄAAGTAC	AACATAGGCG	GATACGAAAA	.CC 280
CCGTATCTT	ACGCGATGT	TTACAGGCTC	TTCTCTCGCA	AC 320
GGGTTCAAC	AGIGCAGGC	GATAGAGGAA	SEAKATITII!	GA 360
AATGTAGCT	TTCAATCCC	GCGGGAGGIA	ATCCACCACCC	TT 400
4	10	420	430	440
	سيليينا	سلسلن	سلسلب	4
TŢĄAĄĄĢCĄ	GGGCAAACG	GCTTTTGCTA	CATAAACGAC	CC 440
CGCTGTGGG	AATTGAGTA	CTTGAGAAAZ	AAAGGCTTTA	AG 480
AGAATACTC	TACATAGAC	CITGATGCCC	CACCACTGCGA	CG 520
GIGITCAGG	AAGCCTTTT	ACGATACAGA	ACCAGGIGIIC	GT 560
CCTGTCCCT	TCACCAGIC	GCCCGAGTAC	GCCTTTCCCT	TT 600
6	10	620	630	640
_			630 	
تسلست	Lullu	سلنسك		سلّ
GAGAAGGGC	TTCCTGGAG	GAGATAGGAG	سلسلك	AA 640
GAGAAGGGC AGGGCTACA	TTCCTGGAO ACCTGAACA	GAGATAGGAG	GAAGGAAAAGG	AA 640 AA 680
GAGAAGGCC AGGCCTACA CGACAACGA ATAGTCAAA	TTCCTGGAG ACCTGAACA GTTCCTCTT GAAGTATTT	GAGATAGGAG TTCCCCIGCC TGCCCTAGAA GAGCCCGAGG	AAAGGAAAAGG AAAGGGCITG AAAGGCITCI ATTTACCITCI	AA 640 AA 680 AA 720 TC 760
GAGAAGGCC AGGCCTACA CGACAACGA ATAGTCAAA	TTCCTGGAG ACCTGAACA GTTCCTCTT GAAGTATTT	GAGATAGGAG TTCCCCIGCC TGCCCTAGAA GAGCCCGAGG	CAAGGAAAAGG CAAAGGGCTTG CAAAGTCTCTGG	AA 640 AA 680 AA 720 TC 760
GAGAAGGCC AGGCCTACA CGACAACGA ATAGTCAAA AACTCGGAA	TTCCTGGAG ACCTGAACA GTTCCTCTT GAAGTATTT CTGACCCAC	GAGATAGGAG TTCCCCIGCC TGCCCTAGAA GAGCCCGAGG	AAAGGAAAAGG AAAGGGCITG AAAGGCITCI ATTTACCITCI	AA 640 AA 680 AA 720 TC 760
GAGAAGGCC AGGCCTACA CGACAACGA ATAGTCAAA AACTCGGAA	TTCCTGGAG ACCTGAACA' GTTCCTCTT GAAGTATTT CTGACCCAC	GAGATAGGAG TTCCCCTGCC TGCCCTAGAA GAGCCCGAGG	SAAGGAAAAGG CAAAGGGCITG AAAACCICIGG SITTACCITCI ATTACCTITCC	AA 640 AA 680 AA 720 TC 760 AA 800
GAGAAGGCC AGGCTACA CGACAACGA ATAGTCAAA AACTCGGAA	TTCCTGGAG ACCTGAACA' GTTCCTCTT GAAGTATTT CTGACCCAC	GAGATAGGAG TTCCCCTGCC TGCCCTAGAA GAGCCCGAGG TCCTTGAAGA 820	AAAGGAAAAGG AAAAGGGCITG AAAAGCTCICG ATTACCTTCI ATTACCTTTCC	AA 640 AA 680 AA 720 TC 760 AA 800
GAGAAGGCC AGGCCTACA CGACAACGA ATAGTCAAAA AACTCGGAA	TTCCTGGAG ACCTGAACA' GTTCCTCTT GAAGTATTT CTGACCCAC	GAGATAGGAG TTCCCCTGCC TGCCCTAGAA GAGCCCGAGG TCCTTGAAGA 820 LLLLLLLLLLLLLLLLLLLLLLLLLLLLLLLLLLL	SAAGGAAAAGG CAAAGGGCTTG AAAA'CTCTGG STTTACCTTCT ATTACCTTTCC	AA 640 AA 680 AA 720 TC 760 AA 800 840 AC 840
GAGAAGGGC AGGGCTACA CGACAACGAA ATAGTCAAAA AACTCGGAAA GTTCAACCTA ATCGTTCGT	TTCCTGGAG ACCTGAACA GTTCCTCTT GAAGTATTT CTGACCCAC 10 LLLLL CTCAAACGT GAGGTTTTC	GAGATAGGAG TTCCCCTGCC TGCCCTAGAA GAGCCCGAGG TCCTTGAAGA 820 LLLLLL TGCCTTTTTA	EAAGGAAAAGG EAAAGGGCITG AAAATCICIGG EITTACCITCI AITTACCITTCC 830 AAAAGCITTCA	AA 640 AA 680 AA 720 TC 760 AA 800 AA 800 AA 840 AC 840 AAC 840
GAGAAGGCC AGGCTIACA AGGCCTIACA ATAGTCAAAA AACTCGGAA BTCAACCTC ATCGTTCGTC GAGGCCGGAT	TTCCTGGAG ACCTGAACA GTTCCTCTT GAAGTATTT CTGACCCAC 10 LL CTCAAACGT GAGGTTTTC ACCATCCTT	GAGATAGGAG TTCCCCTGCC TGCCCTAGAA GAGCCCGAGG TCCTTGAAGA 820 LLLLLL TGCCTTTTTA GGGGAGGGAG	AAAGGAAAAGGAAAAGGAAAAGCTTCTGGATTACCTTCTCTCTC	AA 640 AA 680 AA 720 TC 760 AA 800 B40 AC 840 AC 840 AG 880 AC 920
GAGAAGGGC AGGGCTACA AGGGCTACA ATAGTCAAAA AACTCGGAA GTTCAACCT ATCGTTCGT GAGGCCGAT CCTAATCTG	TTCCTGGAG ACCTGAACA GTTCCTCTT GAAGTATTT CTGACCCAC 10 LLLLLL CTCAAACGT GAGGTTTTC ACCATCCTT GTGCGAGCT	LLLLLL GAGATAGGAG TTCCCCTGGG TGCCCTAGAG GAGCCCGAGG TCCTTGAAGA 820 LLLLLLL TGCCTTTTTA GGGGAGGGAG ACGCCCTCGC	EAAGGAAAAGG EAAAGGGCTTG EAAAGCTTCTGG ETTACCTTCC 830 EAAAGCTTCA ETATACCTCGG	AA 640 AA 680 AA 720 TC 760 AA 800 B40 AC 840 AC 840 AC 840 AC 920 AA 960
GAGAAGGCC AGGCCTACA AGGCCTACA ATAGTCAAAA AACTCGGAA GTTCAACCT ATCGTTCGTC GAGGCCGATT CCTAATCTG AAGCTAAACC	TTCCTGGAG ACCTGAACA' GTTCCTCTT GAAGTATTT CTGACCCAC 10 LILLI CTCAAACGT GAGGTTTTC GAGGTTTTC ACCATCCTT GTGCGAGCT AATAAAGCA	GAGATAGGAG TTCCCCTGCC TGCCCTAGAA GAGCCCGAGG TCCTTGAAGA 820 LLLLLL TGCCTTTTTA GGGGAGGGAG ACGCCCTCGC AAAGAGCCTTT	EAAGGAAAAGG EAAAGGGCTTG AAAATCTCTGG ETTACCTTTCT ATTACCTTTCC 830 LLLLLLLLLLLLLLLLLLLLLLLLLLLLLLLLLL	AA 640 AA 680 AA 720 TC 760 AA 800 B40 AC 840 AC 840 AC 840 AC 920 AA 960
GAGAAGGCC AGGCCTACA AGGCCTACA ATAGTCAAAA AACTCGGAA GTTCAACCT ATCGTTCGTC GAGGCCGATT CCTAATCTG AAGCTAAACC	TTCCTGGAG ACCTGAACA GTTCCTCTT GAAGTATTT CTGACCCAC 10 LL CTCAAACGT GAGGTTTTC ACCATCCTT GTGCGAGCT AATAAAGCA	GAGATAGGAG TTCCCCTGCC TGCCCTAGAA GAGCCCGAGG TCCTTGAAGA 820 LLLLLL TGCCTTTTA GGGGAGGGAG ACGCCCTCGC AAAGAGCTTT	SAAGGAAAAGG SAAAGGGCTTG SAAAGCTTTCC STTACCTTCC 830 SAAAGCTTTCA STATACCTCGG STATACCTCGG STATACCTCGG STATACCTCGG STATACCTCGG STATACCTCGG STATACCTCGG STAAAGAGTAT 1030	AA 640 AA 680 AA 720 TC 760 AA 800 840 AL AC 840 AC 840 AC 920 AA 960 AA 960 AA 960
GAGAAGGCC AGGCTIACA AGGCTIACA ATAGTCAAAA AACTCGGAA GTTCAACCTA ATCGTTCGTA GAGGCGGAT CCTAATCTG AAGCTAAAC 10	TTCCTGGAG ACCTGAACA GTTCCTCTT GAAGTATTT CTGACCCAC 10 LL CTCAAACGT GAGGTTTTC ACCATCCTT GTGCGAGCT AATAAAGCA	GAGATAGGAG TTCCCCTGCC TGCCCTAGAA GAGCCCGAGG TCCTTGAAGA 820 LLLLLL TGCCTTTTTA GGGGAGGGAG ACGCCCTCGC TTCGGGAAGG TCGGGAAGGTTT .020	SAAGGAAAAGG SAAAGGGCTTG SAAAGCTTTCC STTACCTTCC 830 SAAAGCTTTCA STATACCTCGG STATACCTCGG STATACCTCGG STATACCTCGG STATACCTCGG STATACCTCGG STATACCTCGG STAAAGAGTAT 1030	AA 640 AA 680 AA 720 TC 760 AA 800 840 AL AC 840 AG 880 AG 920 AG 1000 1040 AL
GAGAAGGCC AGCGCTACA CGACAACGA ATAGTCAAAA AACTCGGAA GTTCAACCT ATCGTTCGT CGAGGCCGATT CCTAATCTG AAGCTAAAC 10 ACTTGAAGCT ACTTGAAGCT	TTCCTGGAG ACCTGAACA GTTCCTCTT GAAGTATTT CTGACCCAC 10 LLLLLL CTCAAACGT GAGGTTTCCTAACGTT GAGGTTTCCTAAACGTT ACCATCCTT ATTAAAGCA 10 1 LLLLL AGTTTGACG	GAGATAGGAG TTCCCCTGCC TGCCCTAGAF GAGCCCGAGC TCCTTGAAGF 820 LLLLLL TGCCTTTTTF GGGGAGGGAG AAAGAGCTTT .020 LLLLLL ACGAGGTGGF	EAAGGAAAAGG EAAAGGGCTTG AAATCTCTGG ETTACCTTCT 830 AAAGCTTTCA FTATACCTCGG ETATACCTCGG ETATACCTCGG ETATACCTCGG ETATACCTCGG ETAAAGAGTAT 1030	AA 640 AA 680 AA 720 TC 760 AA 800 840 AL AC 840 AG 880 AC 920 AA 960 AA 960 AA 1000 1040 AL AT 1040
CAGAAGGCC AGGCTACA CGACAACGA ATAGTCAAA AACTCGGAA GTTCAACCT ATCGTTCGT CGAGGCGGAT CCTAATCTG AAGCTAAAC 10 ACTTTGAAGG GCTCGAAACC	TTCCTGGAG ACCTGAACA GTTCCTCTT GAAGTATTT CTGACCCAC 10 LL CTCAAACGT GAGGTTTTC ACCATCCTT AATAAAGCA 110 LL AGTTTGACGA CCTAAAGGA	GAGATAGGAG TTCCCCTGCC TGCCCTAGAA TGCCCTGAAGA GAGCCCGAGG R20 LLLLLL GCCTTTTTA CGCGAGGGAG ACCCCTCCC TTCGGGAAGG AAAGAGCTTT .020 LLLLLL ACGAGGTGGA ACCCCTGGAGGAGGAGGAGGAGGAGGAGGAGGTGGAGGAGGTGGAGGA	SAAGGAAAAGG CAAAGGGCTTG AAATCTCTGG STTACCTTCT ATTACCTTCC 830 AAAAGCTTTCA STATACCTCGG CAAGGGCATCG CAAGGGCATCGG TTAAAGAGTAT 1030 ACCGCTCGTAC	AA 640 AA 680 AA 720 TC 760 AA 800 840 AL AC 840 AG 880 AG 920 AA 960 AG 1000 1040 AT 1040 AT 1040

Figure 15

	10	20	30	40	
سلسب	ليسلير	ستلتب	ببسلست	لببيل	
MKKVKLI	GTLDYGKY	RYPKNHPLK	IPRVSLLLRF	LDAMNL	4 0
IDEKELI	KSRPATKE	ELLEHTEL	YINTLMEAER	SQSVPK	80
GAREKYN	IGGYENPV:	SYAMFTGSS	LATGSTVQAI	EEFLKG	120
NVAFNPA	GCMHHAFK!	SRANGFCYI	NDPAVGIEYL	RKKGFK	160
RILYIDL	DAHHCDGV(QEAFYDIDQ	VFVLSLHQSE	EYAFPF	200
•					
	. 210	220	230	240)
سليثت			230)
	للشلك	سلس		لسل	
EKGFLEE	IGEGKGKG	YNLNIPLPK	ستلسب	LEKSLE	240
EKGFLEE	IGEGKGKG PEVYLLQL	YNLNIPLPK	GLNDNEFLFA	LLLL LEKSLE FLKAFN	240
EKGFLEE IVKEVFE IVREVFG	EGVYLGGG EGVYLGGG	SYHPYALAF STDPLLEDY	GLNDNEFLFA TSKFNLSNVA	LLLL LEKSLE FLKAFN GREVPE	240 280 320

PCT/US00/24700

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Figure 16-1

				Residue	# X	Y	Z	occ.	ъ	
TON		CB :	ALA		45.336	36.880	75.042		B 59.90	Segment ID
ATCM	: 2			2 2		38.631	73.628		52.57	AAAA
ATOM	<u>.</u>		مينة		46.410					àààà ;
ATOM	3		ALA.	2	45.780	39.595	74.052		62.46	AAAA '
ATOM	4		LА	2	47.540	37.826	75.673		58.52	AAAA
ATOM	5		:LA	2	46.568	37.432	74.527	1.00	57.32	AAAA !
ATOM	6		LYS	3	46.890	38.570	72.389		39:51	AAAA
STOM	7	CA :	_YS	3	46.687	39.669	71.440	1.00	29.58	AAAA
ATOM	8	CB	LYS	3	47.855	39.763	73.459	1.00	36.03	AAAA ·
ATOM	9		LYS	. 3	49.217	40.007	71.102	1.00	55.16	AAAA
ATOM	10		_75	3	50.315	40.000	73.039	1.00	65.28	AAAA
ATOM	11		LYS	3.	51.700	40.163	73.655		73.41	AAAA
ATOM	. 12		175	3	52.791	40.047	59.642		69.64	AAAA
ATOM	13		LYS	3	45.407	39.422	70.642		23.29	AAAA
	14		LYS	š	44.984	38.282	73.487		27.41	-AAAA' .
—ATOM				4	44.814	40.498	70.138		25.18	AAAA
HOM.	15		VAL							
ATOM	16		VAL	4	43.585	40.418	69.349		22.20	AAAA ¦
ATOM	17		VAL	4	42.501	41.365	59.887		31.46	AAAA
ATOM	18	CG1		4	41.214	41.202	59.066		26.55	AAAA
ATOM	19	CG2		4	42.244	41.080	71.348		34.98	AAAA
ATOM	20		$\Lambda Y\Gamma$	4	43.983	40.851	67.961		25.33	AAAA
ATOM	21	0	VAL	4	44.557	41.927	67.778		21.19	AAAA
MOTA	22	N	LYS	5	43.554	40.023	56.978		21.32	AAAA
ATOM	. 23	CA	LYS	5	44.052	40.291	55.607	1.00	20.10	AAAA
MOTA	24	CB	LYS	5	45.047	39.214	55.177	1.00	23.35	AAAA
ATOM	25	cg	LYS	5	46.301	39.092	55.049	1.00	23.75	AAAA
MOTA	26		LYS	5	47.183	40.334	£5.919	1.00	23.70	AAAA
ATOM	.27		LYS	5	48.510	40.151	55.669	1.00	24.34	AAAA
ATOM	28		LïS	5	49.351	41.387	66.585		22.04	AAAA
MOTA	29		LYS	5	42.914	40.294	£4.596		20.27	AAAA
ATOM	30		LYS	5	41.949	39.535	54.728		13.48	AAAA
ATOM .	31		LEU	6	43.071	41.111	53.564		19.28	AAAA
ATOM	32		LEU	6	42.097	41.156	52.483		20.68	AAAA
	33		LEU	6	41.571	42.574	62.291		23.51	AAAA
ATOM		CG	LEU	6	40.373	42.712	51.342		30.59	AAAA
ATOM	34	CD1		6	40.079	44.192	51.153		29.90	AAAA
ATOM	35			5	40.557	42.085	59.995		38.98	
ATOM	36	CE2.		6	42.964	40.701	51.237		13.17	AAAA
ATOM	37	C	LEU	. б.			50.919		22.31	AAAA
ATOM	38		LEU	. 6 .	43.911	41.249			19.15	AAAA
ATOM	39	N.	ILE		42.359	39.689	50.538		13.38	AAAA
MOTA	40	CA	ILE	7	43.945	39.199	59.338			
ATOM	41	CE	ILE	7	42.922	37.674	59.191		19.05	: 5655
ATOM	42		TLE	7	43.930	37.162	53.144		16.45	ÁÁÁÁ
ATOM	÷ 3		ILE	7	43.253	37.007	60.521		22.91	AAAA
ATOM	14	CD1	ILE	7	43.296	35.543	50.450		34.99	AAAA
ATOM	45	C	ILE	7	42.396	39.850	58.125	1.00	17.95	AAAA .
MOTA	46	0	ILE	7	41.188	39.729	57.928		19.07	AAAA
ATOM	47	N	GLY	9	43.193	40.562	57.330		17.70	AAAA
ATOM	48	·CA	GLY	9	42.623	41.193	55.148		13.11	AAAA
ATOM	49	C	GLY	3	43.640	41.957	55.243		20.91	AAAA
ATOM	50	O	GLY	8	44.849	41.840	55.504	1.00	22.27	AAAA
ATOM	51	N	THR	9	43.134	42.428	54.155		23.99	AAA.
ATOM	52	CA	THR	9	43.950	43.141	53.183	1.00	25.95	AAAA ,
ATOM	53	CE	THR	9	44.739	42.195	52.263	1.00	25.80	AAAA
ATOM	54		THR	9	45.321	42.962	51.199	1.00	25.56	AAAA
ATOM	55		THR	9	43.823	41.144	51.657	1.00	25.24	AAAA
ATOM	56	c	THR	9	43.025	43.957	52.294		29.04	AAAA !
	57	ò	THR	9	41.872	43.582	52.082	1.00	23.05	AAAA
ATOM	58	N	LEU	10	43.517	45.079	51.781		29.19	AAAA
ATOM		CA.	LEU	10	42.690	45.896	50.895		32.55	AAAA
ATCM	59		LEU	10	43.256	47.319	50.761		28.09	AAAA
ATCM	60	CB		10	43.142	48.256	51.958		33.00	AAAA
ATOM	61	CG	LEU				52.347		26.65	AAAA
ATCM	,62	CDI	LEU	10	÷1.580	48.403	53.126		41.33	AAAA
ATCM	63		LEU	10	43.938	47.744	49.512	1.00	32.68	AAAA
ATOM	64	С	LEU	10	42.566	45.261				AAAA
ATCM	65	0	LEU	10	41.736	45.684	48.702		26.97	
ATOM	5 จิ	N	ASP	11	43.377	44.234	;9.25 6	1.00	25.75	AAAA

Figure 16-2

MOTA	67 CA ASP		43.36	7 43.543	47.970	1.00 35.74	AAAA
ATOM	68 CB ASP	11	44.47	7 42.485	47.922		AAAA
MOTA	69 CG ASP	11	45.858		48.079	1.00 46.75	AAAA
MOTA	70 OD1 ASP	11	46.110	44.136			AAAA
MOTA	71 OD2 ASP	11	46.690	42.528	3 43.821		AAAA
MOTA	72 C ASP	11	42.034		47.607		AAAA
MOTA	73 O ASP	11	41.748	42.696	46,420		AAAA
MOTA	74 N TYR	12	41.220	42.558			AAAA
MOTA	75 CA TYR	12	39.923				AAAA
MOTA	76 CB TYR	12	39.119	41.720	49.601		AAAA
MOTA	77 CG TYR	12	39.648				AAAA
MOTA	78 CD1 TYR	12	40.137	40.846	51.755		AAAA
MOTA	79 CE1 TYR	12	40.592	39.808	52.572	1.00.30.35	AAAA
MOTA	80 CD2 TYR	12	39.629			1.00 22.97	AAAA
ATOM	81 CE2 TYR	12	40.077	38.228	50.822	1.00 19.60	AAAA
ATOM	82 CZ TYR	12	40.554	38.499	52.096	1.00 21.42	AAAA
MOTA	83 OH TYR	12	40.964	37.456	52.907	1.00 23.49	AAAA
ATOM	84 C TYR	12	39.144	42.907	47.390	1.00 26.67	AAAA
ATOM	85 O TYR	12	38.307	42.466		1.00 30.51	AAAA
ATOM	86 N GLY	13	39.441		47.492	1.00 30.22	AAAA
MOTA	87 CA GLY	13	. 38.767		46.675	1.00 25.13	AAAA
ATOM	88 C GLY	13	38.911	45.009	45.177	1.00 27.31	AAAA
ATOM	89 O GLY	13	38.096	45.522		1.00 29.38	AAAA
ATOM	90 N LYS	14	39.937	44.269	44.755	1.00 33.56	AAAA
ATOM	91 CA LYS	14	40.176	44.005	43.337	1.00 39.81	AAAA
MOTA	92 CB LYS 93 CG LYS	14	41.680	44.026	43.031	1.00 51.10	AAAA
MOTA MOTA		14		45.424	42.907	1.00 64.99	AAAA
MOTA		14	41.757	46.218	41.692	1.00 72.74	AAAA
ATOM	95 CE LYS 96 NZ LYS	14	42.183	45.639	40.336	1.00 67.25	AAAA
ATOM	97 C. LYS	14	41.637		40.045	1.00 70.06	AAAA
ATOM	98 O LYS	14 14	39.589	42.688	42.834	1.00 39.98	AAAA
ATOM	99 N TYR	15	39.746 38.927	42.350	41.658	1.00 46.99	AAAA
ATOM	100 CA TYR	15	38.318	41.944	43.717	1.00 32.64	AAAA
ATOM	101 CB TYR	15	38.996	40.655 39.512	43.355	1.00 41.01	AAAA
ATOM	102 CG TYR	15	40.496	39.571	44.126	1.00 26.48	AAAA
ATOM	103 CD1 TYR	15	41.289	39.401	44.033 45.167	1.00 34.97 1.00 43.28	AAAA
ATOM	104 CE1 TYR	15	42.677	39.548	45.106	1.00 45.28	AAAA
MOTA	105 CD2 TYR	15	41.127	39.879	42.827	1.00 40.78	AAAA
ATOM	106 CE2 TYR	15	42.508	40.027	42.756	1.00 37.13	AAAA AAAA
ATOM	107 CZ TYR	15	43.275	39.865	43.899	1.00 36.87	AAAA
ATOM	108 OH TYR	15	44.644	40.044	43.844	1.00 35.40	AAAA
ATOM	109 C TYR	15	36.838	40.705	43.714	1.00 38.62	AAAA
ATOM	110 O TYR	15	36.344	39.868	44.468	1.00 37.82	AAAA
MOTA	111 N ARG	16	36.141	41.703	43.177	1.00 44.85	AAAA
MOTA	112 CA ARG	16	34.716	41.890	43.431	1.00 45.75	AAAA
MOTA	113 CB ARG	16	34.320	43.348	43.187	1.00 54.17	AAAA
MOTA	114 CG ARG	16	35.170	44.399	43.875	1.00 66.77	AAAA
MOTA	115 CD ARG	16	34.920	44.506	45.369	1.00 72.39.	AAAA
MOTA	116 NE ARG	16	35.649	45.646	45.923	1.00 85.39	AAAA
MOTA	117 CZ ARG	16	35.489	46.906		1.00 81.94	AAAA
MOTA	118 NH1 ARG	16	34.624	47.197	44.554	1.00 80.19	AAAA
ATOM	119 NH2 ARG	16	36.205	47.878	46.069	1.00 85.46	AAAA
ATOM	120 C ARG	16	33.915	41.029	42.460	1.00 43.50	AAAA
ATOM	121 O ARG	16	34.400	40.667	41.385	1.00 38.62	AAAA
ATOM	122 N TYR	17	32.689	40.692	42.833	1.00 32.68	AAAA
ATOM	123 CA TYR	17	31.850	39.923	41.930	1.00 37.55	AAAA
ATOM	124 CB TYR	17	30.662	39.306	42.672	1.00 41.05	AAAA
ATOM	125 CG TYR	17	31.040	38.104	43.519	1.00 37.51	AAAA
ATOM	126 CD1 TYR	17	32.039	38.194	44.493	1.00 32.59	AAAA
MOTA	127 CE1 TYR	17	32.383	37.095	45.277	1.00 29.32	AAAA
ATOM	128 CD2 TYR	17	30.393	36.875	43.346	1.00 31.46	AAAA
MOTA	129 CE2 TYR	17	30.726	35.772	44.122	1.00 28.64	AAAA
ATOM	130 CZ TYR	17	31.721	35.887	45.088	1.00 27.14	AAAA
MOTA	131 OH TYR	17	32.044	34.807		1.00 21.73	AAAA
ATOM	132 C TYR	17	31.380	40.871	40.836	1.00 40.97	AAAA

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	122	0	WAD.	. 17	71 475	42 007	40 004	1 00 20 50	
ATCM	133	0	TYR		31.435	42.097	40.984	1.00 29.58	AAAA
ATOM	134	N	PRO	18	30.904		39.722	1.00 41.02	AAAA
MOTA	135	CD	PRO	√ 18	30.760	38.910	39.318	1.00 48.67	AAAA
ATCM	136	CA	PRO	18	30.459	41.197	38.649	1.00 49.35	AAAA
ATCM	137	CB	PRO	18	30.321	40.228	37.481	1.00 59.04	AAAA
ATCM	138	CG	PRO	18	29.756	39:017	38.179	1.00 54.15	AAAA
	139						38.864		
ATOM		-	PRO	18	29.178	41.982		1.00 54.97	AAAA
MOTA	140	0	PRO	18	28.457	41.823	39.850	1.00 46.85	AAAA
ATOM	141	N	LYS	. 19	28.961	42.868	37.904	1.00 60.87	AAAA
ATOM	142	CA	LYS	19	27.777	43.696	37.749	1.00 67.78	AAAA
ATOM	143	CB	LYS	19	- 27.155	43.278	36.425	1.00 73.26	AAAA
ATOM	144	ĊG	LYS	19	26.971	41.752	36.414	1.00 77.87	AAAA
	145	CD	LYS	19		41.166	35.209		
MOTA					26.276			1.00 81.01	AAAA
ATOM	146	CE	LYS	19	26.039	39.680	35.471	1.00 82.45	AAAA
MOTA	147	NZ	LYS	19	25.417	38.959	34.331	1.00 83.11	AAAA
MOTA	148	C .	LYS	19	26.688	43.594	38.814-	1.00 64.15	AAAA
ATOM	149	0	LYS	19	26.810	44.047	39.949	1.00 65.73	AAAA
ATOM	150	N	ASN	20	25.604	42.986	38.345	1.00 59.78	AAAA
ATOM	151	CA	ASN	20	24.353	42.703	39.025	1.00 59.91	AAAA
	152	CB	ASN	20	23.516	41.844	38.077	1.00 68.08	
MOTA									AAAA
MOTA	153	CG	ASN	. 20	22.108	42.355	37.907	1.00 78.73	AAAA
MOTA	154		. ASN	20.	21.894	43.498	37.496	1.00 78.67	AAAA
MOTA	155	ND2	ASN	20	21.132	41.505	38.211	1.00 83.22	AAAA
MOTA	156	С	ASN	20	24.474	41.977	40.361	1.00 53.35	AAAA
ATOM	157	0	ASN	20	23.611	42.112	41.234	1.00 59.92	AAAA
ATOM	158	N	HIS	21	25.543	41.206	40.511	1.00 44.23	AAAA
	159	CA	HIS	21	25.768	40.397	41.707	1.00 28.15	AAAA
ATCM									
MOTA	160	CB	HIS	21	27.088	39.639	41.570	1.00 31.84	AAAA
MOTA	161	CG	HIS	. 21	27.155	38.411	42.418	1.00 34.79	AAAA
ATOM T	162		HIS	21	27.344	38.259	43.752	1.00 25.03	AAAA
ATOM	163	ND1	HIS	21	26.929	37.148	41.917	1.00 34.81	AAAA
ATOM	164	CE1	HIS	21	26.979	36.269	42.900	1.00 17.01	AAAA
ATOM	165	NE2	HIS	21	27.228	36.917	44.026	1.00 32.31	AAAA
ATOM	166	С	HIS	21		-41.135	43.051	1.00 29.37	AAAA
	167	õ	HIS	21	26.346	42.210	43.186	1.00 28.54	AAAA
ATOM						40.565	44.066		
ATOM	168	N	PRO	22	25.093			1.00 29.14	AAAA
ATOM	169	CD	PRO	22	24.301	39.322	44.061	1.00 31.20	AAAA
ATOM	170	CA	PRO	22	25.034	41.185	45.395	1.00 32.84	AAAA
ATOM	171	CB	PRO	22	24.174	40.192	46.187	1.00 34.98	AAAA
ATOM	172	CG	PRO	22	23.257	39.634	45.109	1.00 30.11	AAAA
ATOM	173	C	PRO	22	26.411	41.415	46.044	1.00 34.37	AAAA
ATOM	174	O	PRO	22 -	26.554	42.272	46.916	1.00 29.17	AAAA
	175	N	LEU	23	27.415	40.644	45.629	1.00 29.22	AAAA
ATOM			LEU	23	28.765	40.781		1.00 26.49	AAAA
ATOM	176	CA					46.181		
ATOM	177	CB	LEU	23	29.414	39.397	46.332	1.00 22.30	AAAA
ATOM	178	CG	LEU	23	28.703	38.527	47.380	1.00 21.04	AAAA
ATOM	179	CD1	LEU	23	29.307	37.113	47.410	1.00 19.35	AAAA
ATOM	180	CD2	LEU	23	28.850	39.197	48.746	1.00 26.51	AAAA
ATOM	181	С	LEU	23	29.661	41.718	45.361	1.00 25.81	AAAA
ATOM	182	0	LEU	23	30.893	41.693	45.477	1.00 28.45	AAAA
	183	N	LYS	.24	29.018	42.539	44.532	1.00 24.86	AAAA
ATOM	184	CA	LYS	24	29.696	43.552	43.723	1.00 27.35	AAAA
ATOM	185	CB	LYS	24	28.662	44.244	42.830	1.00 28.57	AAAA
ATOM	186	CG	LYS	24	29.118	45.532	42.171	1.00 52.95	AAAA
ATOM	187	CD	LYS	24	28.025	46.603	42.283	1.00 63.74	AAAA
ATOM	188	CE	LYS	24	26.688	46.138	41.706	1.00 66.09	AAAA
	189	NZ	LYS	24	25.595	47.137	41.896	1.00 66.00	AAAA
MOTA	•	C	LYS	24	30.332	44.592	44.676	1.00 29.52	AAAA
ATOM	190								AAAA
ATOM	191	0	LYS	24	31.412	45.123	44.420	1.00 30.67	
ATOM	192	N	ILE	25	29.652	44.879	45.779	1.00 26.90	AAAA
ATOM	193	CA	ILE	25 '	30.151	45.865	46.738	1.00 25.02	AAAA
ATOM	194	CB	ILE	25	29.105	46.177	47.824	1.00 28.34	AAAA
ATOM	195	CG2		25	27.961	46.951	47.237	1.00 23.84	AAAA
	196	CG1		25	28.661	44.869	48.495	1.00 30.31	AAAA
ATOM		CD1		25	27.718	45.051	49.660	1.00 44.90	AAAA
ATOM	197			25				1.00 32.19	AAAA
ATOM	198	С	İFE	وع	31.424	45.463	47.483	1.00 34.13	nnnn

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		_							
MOTA	199		ILE		31.736			1.00 26.54	aaaa
ATOM	200	N	PRO	26	32.191	46.463	47.956	1.00 30.14	AAAA
ATOM	201	CI) PRO	26	31.979	47.907	47.770	1.00 36.38	AAAA
ATOM	202				33.431			1.00 30.63	
									AAAA
MOTA	203				34.014			1.00 34.29	AAAA
ATOM -	204	CG	PRO	26	33.397	48.373	47.617	1.00 43.39	AAAA
ATOM	205	С	PRO	26	32.943	45.727	50.061	1.00 25.99	AAAA
	206		PRO						•
MOTA					31.854	46.110		1.00 25.51	AAAA
MOTA	. 207	N	ARG	27	33.719	44.880	50.743	1.00 21.98	AAAA
ATOM	208	CA	ARG	27	33.267	44.347	52.035	1.00 26.17	AAAA
ATOM	209	CE	ARG	27	32.641	42,969	51.834	1.00 22.70	
		•							AAAA
ATOM	210			. 27	31.442	43.039	-	1.00 26.75	AAAA
ATOM	211	CD	ARG	27	30.832	.41.672	50.581	1.00 33.22	AAAA
ATOM	212	NE	ARG	27	30.121	41.098	51.716	1.00 28.66	AAAA
ATOM	213	CZ	ARG	27	30.582	40.129	52.503	1.00 31.79	AAAA
	214		1 ARG	27					
MOTA					31.778	39.598	52.290	1.00 34.08	- AAAA
ATOM	215	NH	2 ARG	27	29.833	39.688	53.505	1.00 26.16	AAAA
ATOM	216	С	ARG	27 ·	34.358	44.297	53.090	1.00 24.10	AAAA
ATOM	217	0	ARG	27	34.326	45.074	54.038	1.00 23.50	
									AAAA
MOTA	218	N	VAL	28	35.314	43.390	52.960	1.00 21.45	AAAA
ATOM	219	CA	VAL	28	36.385	43.385	53.953	1.00 21.75	AAAA
MOTA	220	CB	VAL	28	37.221	42.101	53.866	1.00 26.55	AAAA
ATOM	221	CG	1 VAL	28	38.407	42.177	54.830	1.00 23.84	
									AAAA
MOTA	222		2 VAL	28	36.337	40.906	54.214	1.00 19.20	AAAA
MOTA	223	С	VAL	28	37.277	44.611	53.736	1.00 20.86	AAAA
MOTA	224	0	VAL	28	37.770	45.223	54.702	1.00 25.15	AAAA
ATOM	225	N.	SER	29	37.480	44.996	52:475	1.00 19.22	AAAA
MOTA	226	CA	SER	. 29	. 38.320	46.169	52.209	1.00 19.63	AAAA
MOTA	227	CB	SER	29	38.591	46.352	50.702	1.00 24.45	AAAA
MOTA	228	OG	SER	29	37.411	46.697	49.984	1.00 28.74	AAAA
ATOM	229	С	SER	29	37.579	47.381	52.756	1.00 21.50	AAAA
ATOM	230	ŏ	SER	29	38.184		53.271		
						48.320		1.00 18.95	AAAA
ATOM	231	N	LEU	30	36.256	47.353	52.673	1.00 19.56	AAAA
MOTA	232	CA	LEU	30	35.499	48.481	53.177	1.00 25.97	AAAA
MOTA	233	CB	LEU	30	34.032	48.396	52.744	1.00 22.90	AAAA
ATOM	234	CG	LEU	30	33.085	49.541	53.157	1.00 26.62	AAAA
MOTA	235		LEU	30	32.885	49.539	54.648	1.00 38.27	` AAAA
ATOM	236	CD2	LEU	30	33.653	50.885	52.698	1.00 25.71	AAAA
ATOM	237	С	LEU	30	35.604	48.509	54.696	1.00 18.44	AAAA
ATOM .	238	0	LEU	30	35.704	49.580	55.273	1.00 25.05	AAAA
	239	N	LEU	31					
ATOM					35.578	47.336	55.336	1.00 19.65	AAAA
MOTA	240	CA	LEU	31	35.672	47.270	56.797	1.00 20.47	AAAA
ATOM	241	CB	LEU	31	35.613	45.821	57.300	1.00 20.60	AAAA
ATOM	242	CG	LEU	31	34.988	45.456	58,665	1.00 39.80	AAAA
	243		LEU	31	35.712	44.219	59.257	1.00 23.99	
ATOM									AAAA
MOTA	244	CD2	LEU	31	35.085	46.591	59.637	1.00 28.48	AAAA
ATOM	245	С	LEU	31	37.009	47.870	57.229	1.00 23.85	AAAA
ATOM	246	0	LEU	31	37.070	48.673	58.154	1.00 21.24	AAAA
	247	N	LEU	32	38.079	47.462	56.562	1.00 23.91	
ATOM									AAAA
ATCM	248	CA	LEU	32	39.400	47.965	56.899	1.00 24.82	AAAA
MOTA	249	CB	LEU	32	40.479	47.320	56.018	1.00 24.81	AAAA
ATOM	250	CG	LEU	32	40.849	45.854	56.276	1.00 27.00	AAAA
				32					
ATOM	251		LEU		41.995	45.435	55.354	1.00 27.13	AAAA
MOTA	. 252	CD2	LEU	32	41.285	45.687	57.720	1.00 34.49	AAAA
ATOM	253	С	LEU	32	39.466	49.475	56.763	1.00 19.56	AAAA
ATOM	254	0	LEU	32	39.958	50.143	57.662	1.00 20.71	AAAA
MOTA	255	Vi	ARG	33	38.974	50.006	55.645	1.00 23.25	AAAA
ATOM	256	CA	ARG	33	39.007	51.449	55.441	1.00 24.33	AAAA
ATOM	257	CB	ARG	33	38.575	51.806	54.013	1.00 23.46	AAAA
	258	CG	ARG	33	39.571	51.327	52.945	1.00 26.94	AAAA
MOTA									
ATOM	259	CD	ARG	33	39.337	51.976	51.585	1.00 42.13	AAAA
MOTA	260	NE	ARG	33	38.023	51.661	51.037	1.00 59.06	AAAA
ATOM	261	CZ	ARG	33	37.583	52.088	49.857	1.00 60.87	AAAA
	262	NH1		33	38.353	52.850	49.095	1.00 65.33	AAAA
MOTA									
MOTA	263	NH2		33	36.373	51.743	49.433	1.00 56.24	AAAA
ATOM	264	С	ARG	33	38.124	52.156	56.455	1.00 30.33	AAAA
							-		

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	265 0	200	32	20 441	E2 252	56.905	1 00 25 45	
MOTA	265 0	ARG	33	38.441	53.252		1.00 25.45	AAAA
ATOM	266 N	PHE	34	37.022	51.514	56.828	1.00 24.98	AAAA
	267 CA	PHE	34	36.099	52.085	57.789	1.00 27.09	AAAA
MOTA								
ATOM	268 CB	PHE	34	34.798	51.276	57.807	1.00 24.88	AAAA
MOTA	269 CG	PHE	34	33.719	51.898	58.631	1.00 20.46	AAAA
MOTA	270 CD	1 PHE	34	33.043	53.018	58.171	1.00 18.74	AAAA
MOTA	271 CD2	2 PHE	34	33.396	51.383	59.889	1.00 20.19	AAAA
				_				
MOTA	272 CE	1 PHE	34	32.043	53.627	58.956	1.00 23.04	AAAA
MOTA	273 CE2	2 PHE	· 34.	32.406	51.974	60.681	1.00 25.08	AAAA
		PHE	34	31.726	53.104	60.209	1.00 23.31	
MOTA	274 CZ							АААА
ATOM	275 C	PHE	34	36.709	52.115	59.194	1.00 23.93	AAAA
ATOM	276 O	PHE	34	36.668	57 178	59.883	1:00 21.71	'AAAA
ATOM	277 N	LYS	35	37.298	51.013	59.645	1.00 21.33	AAAA
MOTA	278 CA	LYS	35	37.862	51.084	60.978	1.00 22.54	AAAA
			35		49.716		1:00 29.70	
MOTA	279 CB	LYS		38.276		61.476		AAAA
MOTA	280 CG	LYS	35	37.082	48.890	61.924	1.00 29.48	AAAA
MOTA	281 CD	LYS	35	37.517	47.535	62.398	1.00 42.17	AAAA
MOTA	282 CE	LYS	35	38.157	46.762	61.275	1.00 34.89	AAAA
ATOM	283 NZ	LYS	35	39.372	47.412	60.719	1.00 67.18	AAAA
			35					
ATOM	284 C	LYS		39.027	52.055	61.040	1.00 24.68	AAAA
ATOM	285 O	LYS	35	39.282	52.640	62.085	1.00 22.33	AAAA
ATOM	286 N	ASP	36	39.724	52.231	59.926	1.00 25.67	AAAA
MOTA	287 CA	ASP	36	40.842	53.163	59.898	1.00 25.57	AAAA
ATOM	288 CB	ASP	36	41.669	52.984	58.621	1.00 32.26	AAAA
		ASP	36	42.881	53.914	58.572	1.00 33.92	
MOTA	_							AAAA
MOTA	290 OD1	. ASP	36	43.641	53.969	59.563	1.00 40.22	AAAA
ATOM	291 OD2	ASP	36	43.078	54.575	57.538	1.00 40.06	AAAA
						59.973		
MOTA	292 C	ASP	36	40.285	54.578		1.00 28.04	AAAA
ATOM	293 O	ASP	36	40.761	55.397	60.765	1.00 29.52	AAAA
ATOM	294. N	ALA	37	39.272	54.864	59.159	1.00 23.32	AAAA
ATOM	295 CA	ALA	37	38.651	56.192	59.163	1.00 28.22	AAAA
ATOM	296 CB	ALA	37	37.506	56.251	58.119	1.00 25.93	AAAA
	297 C	ALA	37	38.127	56.549	60.565	1.00 28.41	AAAA
ATOM								
ATOM	298 O	ALA	37	38.186	57.708	60.972	1.00 29.27	AAAA
MOTA	299 N	MET .	38	37.639	55.547	61.300	1.00 24.76	AAAA
	300 CA	MET	38	37.103	55.727	62.669	1.00 25.45	AAAA
ATOM								
ATOM	301 CB	MET	38	36.077	54.625	62.982	1.00 25.19	AAAA
MOTA	302 CG	MET	38	34.816	54.660	62.148	1.00 22.32	AAAA
	303 SD	MET	38	33.733	55.983	62.702	1.00 29.90	AAAA
MOTA							, .	
ATOM	304 CE	MET	38	33.402	55.417	64.376	1.00 26.51	AAAA
ATOM	305 C	MET	38	38.203	55.667	63.744	1.00 26.42	AAAA
			38			64.947	1.00 23.77	AAAA
ATOM	306 O	MET		37.924	55.818			
ATOM	307 N	asn	39	39.437	55.434	63.300	1.00 26.21	AAAA
ATOM	308 CA	ASN	39	40.607	55.308	64.170	1.00 28.53	AAAA
						64.855	1.00 33.95	AAAA
ATOM	309 CB	ASN	39	40.926	56.643			
ATOM	310 CG	ASN	39	41.153	57.751	63. 8 58	1.00 29.46	AAAA
ATOM	311 OD1	ASN	39	41.930	57.596	62.925	1.00 36.28	AAAA
ATOM		ASN	39	40.472	58.880	64.046	1.00 40.03	AAAA
ATOM	313 C	ASN	39	40.374	54.223	65.205	1.00 30.07	AAAA
		ASN	39	40.682	54.390	66.395	1.00 25.47	AAAA
ATOM								
ATOM	315 N	LEU	40	39.814	53.105	54.744	1.00 28.19	AAAA
ATOM	316 CA	LEU	40	39.527	51.984	65.633	1.00 25.50	AAAA
-							1.00 32.14	AAAA
MOTA	317 CB	LEU	40	38.060	51.562	65.514		
MOTA	318 CG	LEU	40	37.044	52.585	66.036	1.00 30.47	AAAA
ATOM	319 CD1	LEU	40	35.637	52.027	65.894	1.00 29.07	AAAA ·
	_							
MOTA		LEU	40	37.325	52.889	67.491	1.00 23.80	AAAA
ATOM	321 C	LEU	40	40.433	50.771	65.415	1.00 26.99	AAAA
	322 0	LEU	40	40.157	49.683	65.915	1.00 25.41	AAAA
ATOM								
ATOM	323 N	ILE	41	41.528	50.970	64.691	1.00 28.33	AAAA
ATOM	324 CA	ILE	41	42.459	49.882	64.459	1.00 25.08	AANA
						63.243	1.00 25.01	AAAA
ATOM	325 CB	ILE	41	42.010	49.020			
MOTA:	326 CG2	ILE	41	42.061	49.824	61.961	1.00 22.74	AAAA
		ILE	41	42.917	47.802	63.128	1.00 31.01	AAAA
atom								
ATOM	328 CD1		41	42.895	46.951	64.341	1.00 42.18	AAAA
ATOM	329 C	ILE	41	43.900	50.376	64.247	1.00 24.09	AAAA
	330 0	ILE	41	44.128	51.406	63.621	1.00 28.92	AAAA
MOTA					3400	-		
	•				•			

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MOTA MOTA	33 33	2 C			44.86 46.27				AAAA
ATOM		_	A ASP	42	46.27				
	22.								2222
MOTA	33:	3 C	B ASP	42		1 . 50.09			AAAA AAAA
MOTA	33	4 C	G ASP	42	46.26				AAAA
ATOM	33!	5 01	DI ASP	42	46.07				AAAA
MOTA	330	o o	D2 ASP	42	45.92				
ATOM	33.	7 C	ASP	42	46.98				AAAA
ATOM	338	3 0	ASP	42	46.59		_		AAAA .
ATOM	339) N	GLU	43	48.03				AAAA
ATOM	340) CZ	4 GLU	43	48.79				AAAA
MOTA	341	L CE	GLU	43	50.078			1.00 36.12	. AAAA
MOTA	342	? C0	GLU	43	49.886				AAAA
ATOM	343	CI	GLU	43	51.214				AAAA AAAA
ATOM	344	OE	:1 GLU	43	51.928				
ATOM	345	OE	2 GLU	43	51.550				AAAA
ATOM	346	C	GLU-	43	49.196				AAAA
MOTA	347	0	GLU	43	49.125				AAAA
MOTA	348	N	LYS	44	49.636				AAAA
MOTA	349	CA	LYS	44	50.084				AAAA
MOTA	350	CB	LYS	44	50.974				AAAA
MOTA	351	CG	LYS	44	52.211			1.00 59.37	AAAA
ATOM	352	CD	LYS	44	53.187			1.00 68.87	AAAA AAAA
MOTA	353	CE	LYS	44	54.373			1.00 67.21	AAAA
MOTA	354	NZ	LYS	44	55.361			1.00 74.00	AAAA
ATOM	355	C	LYS	44	48.982	44.889		1.00 26.75	AAAA
ATOM	356	0	LYS	44	49.265			1.00 27.37	AAAA
MOTA	357	N	GLU	45	47.731	45.278		1.00 29.20	AAAA
MOTA	358	CA	GLU	45	46.580	44.414	65.165	1.00 21.58	AAAA
ATOM	359	CB	GLU	45	45.387	45.243	65.676	1.00 18.24	AAAA
ATOM	360	CG	GLU	45	45.551	45.828	67.077	1.00 26.57	AAAA
ATOM	361	CD	GLU	45	44.418	46.772	67.453	1.00 23.12	AAAA
ATOM	362		LGLU	45	44.224	47.783	66.746	1.00 21.64	AAAA
ATOM	363		GLU	45	43.725	46.509	68.454	1.00 26.48	AAAA
ATOM	364	С	GLU	45 .	46.163	43710	63.870	1.00 26.31	AAAA
ATOM	365	0	GLU	45	45.400	42.739	63.889	1.00 22.32	AAAA
MOTA	366	N	LEU	46	46.674	44.204	62.748	1.00 20.15	AAAA
ATOM	367	CA	LEU	46	46.317	43.642	61.448	1.00 25.80	AAAA
ATOM	368	CB	LEU	46	46.137	44.774	60.433	1.00 27.25	AAAA
ATOM	369	CG	LEU	46	45.763	44.397	58.997	1.00 37.72	AAAA
ATOM	370 371		LEU	46	44.356	43.810	58.984	1.00 39.46	AAAA
ATOM	372		LEU	46	45.822	45.632	58.101	1.00 35.43	AAAA
ATOM ATOM	373	С О	LEU	46	47.305	42.623	60.896	1.00 28.88	AAAA
ATOM	374	N	LEU ILE	46	48.513	42.860	60.862	1.00 31.98	KAAA
ATOM	375	CA		47	46.791	41.469	60.482	1.00 16.92	AAAA
ATOM	376	CB	ILE ILE	47 47	47.638	40.448	59.872	1.00 20.98	· AAAA
ATOM	377		ILE	47	47.412 48.115	39.046	67.513	1.00 21.51	AAAA
ATOM	378		ILE	47	47.947	37.958 39.040	52.696	1.00 20.32	AAAA
ATOM	379			47	49.450	39.207	6 .950	1.00 20.71	AAAA
ATOM	380	C	ILE	47	47.227		62.052	1.00 38.87	AAAA
ATOM	381	ō	ILE	47	46.036	40.417 40.279	58.406 58.101	1.00 24.50 1.00 20.74	AAAA
ATOM	382	N	LYS	48	48.195	40.550	57.500		AAAA
ATOM	383	CA	LYS	48	47.883	40.543	56.072	1.00 18.73	AAAA
ATOM	384	CB	LYS	48	49.095	40.991	55.239	1.00 15.55	AAAA
ATOM	385	CG	LYS	48	48.836	41.011		1.00 16.52	AAAA
ATOM	386	CD	LYS	48	50.072	41.451	53.738 52.957	1.00 23.25	AAAA
ATOM		CE	LYS	18	49.796	41.496	51.462	1.00 32.69 1.00 26.00	AAAA
ATOM		NZ	LYS	48	48.704	42.449			AAAA
ATOM			LYS	18	47.473	39.140	51.114 55.629	1.00 46.33	AAAA
ATOM			LYS	48	48.177	38.174	55.887	1.00 14.43	AAAA
ATCM			SER	19	46.343	39.049	54.945	1.00 16.83	AAAA
ATOM			SER	19	45.838	37.780	54.439	. –	AAAA
ATOM			SER	49	44.517	37.780	53.694	1.00 14.33	. AAAA
ATOM			SER	49	43.509	38.613	54.492	1.00 13.21	AAAA
ATOM			SER	49	46.810	37.131	53.459	1.00 16.86 1.00 24.11	AAAA
ATOM			SER	49	47.463	37.815	52.663	1.00 24.11	AAAA
						- ,		1.00 13.33	AAAA

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Figure 16-7

					46 000	35.805	53.519	1.00 16.83	AAAA
MOTA	397	N	ARG	50	46.890			-	
MOTA	398	CA	ARG	50	47.724	35.037	52.610	1.00 23.88	AAAA
ATOM	399	CB	· ARG	50	48.805	34.247	53.366	1.00 27.48	AAAA
ATOM	400	CG	ARG	50	48.284	33.036	54.177	1.00 22.99	AAAA
ATOM	401	CD	ARG	50	49.453	32.263	54.759	1.00 25.20	AAAA
	402	NE	ARG	50	49.073	31,197	55.684	1.00 15.88	AAAA
ATOM					48.411	30.093	55.368	1.00 14.34	AAAA
MOTA	403	CZ		50	-				AAAA
MOTA	404		ARG	50 '	48.023	29.863	54.117	1.00 15.78	
ATOM	405	NH2	ARG	50	48.150	29.197	56.312	1.00 16.78	AAAA
MOTA	406	С	ARG	50	46.821	34.023	51.905	1.00 20.20	AAAA
ATOM	407	0	ARG	50 -	45.763	33.650	52.414	1.00 18.63	AAAA
	408	N	PRO	51	47.203	33.596	50.699	1.00 15.63	AAAA
ATOM'					48.322	34.028	49.850	1.00 19.45	AAAA
ATOM	409	CD	PRO	51			49.994	1.00 14.35	AAAA
ATOM	410	CA	PRO	51. "		.32:606			
ATOM	411	CB	PRO	51 .	47.076	32.514	48.629	1.00 17.73	AAAA
MOTA	412	CG	PRO	51	47.707	33.890	48.475	1.00 17.62	AAAA
ATOM	413	С	PRO	51	46.452	31.256	50.708	1.00 15.73	AAAA
	414	ō	PRO	51	47.460	30.942	51.350	1.00 18.67	AAAA
ATOM			ALA	52	45.377	30.470	50.618	1.00 11.47	AAAA
MOTA	415	N				29.117	51.161	1.00 9.78	AAAA
MOTA	416	CA	ALA	52	45.375				AAAA
ATOM	417	CB	ALA	52	43.967	28.529	51.112	1.00 12.19	
MOTA	418	С	ALA	52	46.301	28.342	50.209	1.00 17.19	AAAA
ATOM	419	0	ALA	52	46.307	28.609	49.006	1.00 16.46	AAAA
ATOM	420	N	THR	53	47.081	27.392	50.723	1.00 16.40	AAAA
	421	CA	THR	53	47.952	26.615	49.843	1.00 16.32	AAAA
MOTA			THR	53 [.]	49.109	25.959	50.612	1.00 15.82	AAAA
MOTA	422	CB				25.016	51.559	1.00 16.25	AAAA
MOTA	423		THR	53	48.582			1.00 14.34	AAAA
ATOM	424		THR	53	49.923	27.030	51.336		AAAA
ATOM	425	С	THR	53	47.104	25.520	49.215	1.00 14.06	
ATOM	426	0	THR	53	46.012	25.241	49.690	1.00 17.87	AAAA
ATOM	427	N	LYS.	54	47.599	24.903	48.145	1.00 16.10	AAAA
	428	CA	LYS	54	46.848	23.832	47.492	1.00 19.00	AAAA
MOTA	429	CB	LYS	54	47.671	23.245		1.00 22.92	AAAA
ATOM				54	46.955	22.172	45.539	1.00 32.99	AAAA
ATOM	430	.CG	LYS			22.733	44.757	1.00 51.34	AAAA
ATOM	431	CD	LYS	54	45.787			1.00 64.17	AAAA
ATOM	432	CE	LYS	54	46.244	23.565	43.561		
ATOM	433	NZ	LYS	54	46.898	22.733	42.505	1.00 63.45	AAAA
ATOM	434	С	LYS	54	46.554	22.738	48.520	1.00 22.48	AAAA
ATOM	435	0	LYS	54	45.463	22.158	48.555	1.00 19.97	AAAA
	436	N	GLU	55	47.536	22.465	49.364	1.00 25.65	AAAA
ATOM					47.389	21.432	50.383	1.00 25.08	AAAA
MOTA	437	CA	GLU	55			51.116	1.00 25.40	AAAA
MOTA	438	CB	GLU	55	48.718	21.241		1.00 48.95	AAAA
ATOM	439	CG	GLU	55	48.703	20.185	52.199		
ATOM	440	CD	GLU	55	50.106	19.821	52.673	1.00 64.21	AAAA
ATOM	441	OE1	GLU	55	50.220	19.033	53.640	1.00 62.38	AAAA
ATOM	42	OE2	GLU	55	51.093	20.311	52.073	1.00 58.22	AAAA
	.43	c	GLU	55	46.273	21.773	51.362	1.00 18.91	AAAA
ATOM		ō	GLU	55	45.489	20.908	51.723	1.00 17.43	AAAA
ATOM	-44				_	23.029	51.786	1.00 16.80	AAAA
ATOM	445	N	GLU	56	46.196			1.00 17.24	AAAA
MOTA	446	CA	GLU	56	45.137	23.432	52.698		AAAA
ATOM	447	CB	GLU	56	45.399	24.855	53.204	1.00 16.15	
ATOM	448	CG	GLU	56	46.709	24.941	54.009	1.00 14.41	AAAA
ATOM	449	CD	GLU	56	47.087	26.354	54.358	1.00 20.17	AAAA
	450		GLU	56	46.713	27.252·	53.567	1.00 17.12	AAAA
MOTA			GLU	56	47.773	26.564	55.394	1.00 18.23	AAAA
MOTA	451					23.313	52.000	1.00 15.95	AAAA
MOTA	452	C	GLU	56	43.781			1.00 17.82	AAAA
MOTA	453	0	GLU	56	42.799	22.869	52.599		AAAA
MOTA	454	N	LEU	57	43.722	23.691	50.725	1.00 17.53	
MOTA	455	CA	LEU	57	42.466	23.579	49.989	1.00 16.34	AAAA
	456	CB	LEU	57	42.591	24.177	48.586	1.00 13.86	AAAA
ATOM		CG	LEU	57	42.773	25.707	48.552	1.00 15.24	AAAA
ATOM	457			57 57	42.923	26.182	47.101	1.00 19.30	AAAA
ATOM	458		LEU				49.207	1.00 15.14	AAAA
ATOM	459		LEU	57	41.546	26.380			AAAA
ATOM	460	С	LEU	57	42.016	22.126	49.868	1.00 18.46	
ATOM	461	0	LEU	57	40.824	21.823		1.00 17.27	AAAA
	462	N	LEU	58	42.975	21.234	49.636	1.00 16.43	AAAA
ATOM	302				J		-		•
			-						

Figure 16-8

MOTA	463 CA LEU	58	42.662	2 19.822	49:475	1.00 15.18	2222
ATOM	464 CB LEU	58	43.788				AAAA
ATOM	465 CG LEU						AAAA
		58	44.029			1.00 21.72	AAAA
MOTA	466 CD1 LEU	.58	45.221	18.982	46.680	1.00 31.92	AAAA
MOTA	467 CD2 LEU	58	42.786	19.549	46.469		AAAA
MOTA	468 C LEU	58	42.339				
ATOM							AAAA
		58	42.067		50.795	1.00 20.40	AAAA
MOTA	470 N LEU	59	42.377	19.849	51.896	1.00 13.50	AAAA
ATOM	471 CA LEU	59	41.958	19.261			
ATOM	472 CB LEU	59	42.182				AAAA
							AAAA
ATOM		59	43.619			1.00 22.57	AAAA
ATOM	474 CD1 LEU	59	43.640	21.654	55.808	1.00 19.88	AAAA
MOTA	475 CD2 LEU	59 -	44.255	19.253			
ATOM	476 C LEU	59	40.446				AAAA
ATOM	477 O LEU						AAAA
		59	39.897				AAAA
MOTA.	478 N PHE	60 .	39.766		52.179	1.00 14.64	- AAAA
ATOM	479 CA PHE	60	38.338	19.536	51.970	1.00 18.17	AAAA
ATOM	480 CB PHE	60	37.519			1.00 18.80	
ATOM	481 CG PHE	60					AAAA
			36.028			1.00 15.94	AAAA
MOTA	482 CD1 PHE	60	35.320	19.476	52.817	1.00 19.98	AAAA
ATOM	483 CD2 PHE	60	35.339	21.524	51.576	1.00 18.09	AAAA
MOTA	484 CE1 PHE	60	33.947	19.338	52.587	1.00 18.72	
ATOM	485 CE2 PHE	60			52.307	1.00 10.72	AAAA
			33.964	21.399		1.00 19.19	AAAA
ATOM	486 CZ PHE	60	33.268	20.295	51.850	1.00 18.43	AAAA
ATOM	487 C PHE	60	37.916	19.337	50.510	1.00 16.45	AAAA
MOTA	488 O PHE	60	37.227	18.371	50.179	1.00 19.18	AAAA
ATOM	489 N HIS	61	38.308	201257			
ATOM	490 CA HIS	61				1.00 18.26	AAAA
			37.913	20.163	48.235	1.00 14.47	AAAA
ATOM	491 CB HIS	61	38.004	21.545	47.582	1.00 17.15	AAAA
ATOM	492 CG HIS	61	36.968	22.494	48.084	1.00 14.20	AAAA
ATOM	493 CD2 HIS	61	35.645	22.580	47.816	1.00 11.05	AAAA
ATOM	494 ND1 HIS	61	37.237	23.477	49.012		
ATOM	495 CE1 HIS					1.00 23.25	AAAA
		61	36.121	24.131	49.291	1.00 13.35	AAAA
ATOM	496 NE2 HIS	61	35.143	23.606	48.579	1.00 21.07	AAAA
ATOM	497 C HIS	61	38.695	19.157	47.417	1.00 18.29	AAAA
ATOM	498 O HIS	61	39.828	18.819	47.761	1.00 17.50	AAAA
ATOM	499 N THR	62	38.071	18.658			
ATOM	500 CA THR				46.346	1.00 15.39	AAAA
		62	38.741	17.686	45.473	1.00 19.02	AAAA
MOTA	501 CB THR	62	37.734	16.767	44.756	1.00 19.61	AAAA
ATOM	502 OG1 THR	62	36.795	17.548	44.006	1.00 22.05	AAAA
ATOM	503 CG2 THR	62	36.995	15.925	45.767	1.00 28.99	AAAA
ATOM	504 C THR	63		18.398			
					44.440	1.00 23.22	AAAA
ATOM		62	39.311	19.532	44.044	1.00 17.47	AAAA
MOTA	506 N GLU	63	40.657	17.732	44.009	1.00 18.94	AAAA
MOTA	507 CA GLU	63	41.571	18.324	43.046	1.00 22.44	AAAA
ATOM	508 CB GLU	63	42.736	17.384	42.750	1.00 28.31	
ATOM	509 CG GLU	63	43.885	17.476			AAAA
	510 CD GLU				43.708	1.00 60.37	AAAA
MOTA		63	45.154	16.893	43.115	1.00 65.08	AAAA
ATOM	511 OE1 GLU	63	45.603	17.407	42.065	1.00 66.44	AAAA
ATOM	512 OE2 GLU	63	45.697	15.927	43.694	1.00 71.72	AAAA
MOTA	513 C GLU	63	40.983	18.764			
	514 O GLU				41.730	1.00 18.63	AAAA
ATOM		63	41.340	19.827	41.228	1.00 18.37	AAAA
ATOM	515 N ASP	64	40.108	17.943	41.153	1.00 19.77	AAAA
ATOM	516 CA ASP	64	39.508	18.277	39.864	1.00 -17.88	AAAA
MOTA	517 CB ASP	64	38.584	17.159	39.372	1.00 20.43	AAAA
ATOM	518 .CG ASP	64	37.429				
				16.884	40.330	1.00 42.71	AAAA
ATOM	519 OD1 ASP	64	36.415	16.291	39.899	1.00 45.01	AAAA
ATOM	520 OD2 ASP	64	37.537	17.243	41.521	1.00 51.77	AAAA
ATOM	521 C ASP	64	38.701	19.582	39.964	1.00 21.90	AAAA
ATOM	522 O ASP	64	38.726	20.410	39.042		
				20.410		1.00 17.35	AAAA
ATOM		65	37.980	19.750	41.072	1.00 16.17	AAAA
atom	524 CA TYR	65	37.178	20.957	41.292	1.00 15.62	AAAA
ATOM	525 CB TYR	65	36.258	20.796	42.529	1.00 12.04	AAAA
ATOM	526 CG TYR	65	35.501	22.065	42.886	1.00 12.23	
	527 CD1 TYR						AAAA
ATOM		65	34.699	22.718	41.940	1.00 14.73	AAAA
atom	528 CE1 TYR	65	34.028	23.910	42.253	1.00 18.23 -	AAAA
•	•				-		-

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A DOM	529	CD2	TYR	65	35.609	22.631	44.163	1.00 13.67	AAAA
MOTA	-						44.486	1.00 18.16	AAAA
MOTA	530	CEZ	TYR	65	34.943	23.824			
ATOM	531	CZ.	TYR	65.	34.162	24.461	43.533	1.00 16.88	. AAAA
						25.665	43.837	1.00 14.59	λλλλ
MOTA	532	OH	TYR	65	33.555				
MOTA	533	С	TYR	65	38.090	22.177	41.459	1.00 15.27	AAAA
			TYR	65	37.882	23.189	40.798	1.00 15.96	AAAA
ATOM	534	0							
ATOM	535	N	ILE	66	39.098	22.073	42.321	1.00 14.29	AAAA
	536	CA	ILE	66	40.022	23.179	42.540	1.00 18.86	AAAA
MOTA								1.00 15.56	AAAA
MOTA	537	CB	ILE	66	41.090	22.836	43.617		
	538	CG2	ILE	66	42.152	23.943	43.698	1.00 20.45	AAAA
ATOM						22.659	44.967	1.00 19.68	AAAA
MOTA	539	CGI	ILE	66	40.405				
MOTA	540	CD1	ILE	66	39.71 <i>7</i>	23.948	45.454	1.00 29.11	· AAAA
				66	40.716	23.519	41.236	1.00 25.20	AAAA .
MOTA	541	С	ILE						
MOTA	542	0	ILE	66	40.809	24.692	40.895	1.00 14.60	, AAAA
	543	N	ASN	67	41.190	22.508	40.498	1.00 18.21	AAAA
MOTA							39.236	1.00 20.03	AAAA
ATOM	544	CA	ASN	67	41.879	22.789			
ATOM	545	CB	ASN	67	42.448	21.523	38.580	1.00 21.73	AAAA
					•	20.954	39.333	1.00 21.69	AAAA
ATOM	546	CG	ASN	67	43.645				
ATOM	547	OD1	ASN	6 7	44.293	21.645	40.110	1.00 23.97	AAAA
				67	43.947	19.692	39.086	1.00 23.23	AAAA
MOTA	548		ASN						
MOTA	549	С	ASN	67	40.970	23.500	38.250	1.00 15.87	AAAA
	550	0	ASN	67	41.431	24.347	37.473	1.00 18.64	AAAA
ATOM							38.295	1.00 16.55	AAAA
MOTA	551	N	THR	68	39.681	23.180			
MOTA	552	CA	THR	68	38.729	23.814	37.400	1.00 20.34	AAAA
			THR	68	37.360	23.114	37.441	1.00 22.99	AAAA
MOTA	553	CB							
ATOM	554	OG1	THR	68·	37.511	21.760	36.978	1.00 21.75	AAAA
	555		THR	68	36.378	23.827	36.536	1.00 17.37	AAAA
MOTA							37.755		AAAA
ATOM	556	С	THR	68 ·	38.561	25.291			
MOTA	557	0	THR	68	38.472	26.139	36.871	1.00 18.79	AAAA
				69	38.534	25_604	39.045	1.00 14.82	AAAA
ATOM	558	N	LEU						AAAA
MOTA	559	CA	LEU	69	38.405	27.000	39.447	1.00 15.20	
	560	CB	LEU	69	38.295	27.126	40.973	1.00 16.87	AAAA
MOTA						26.551	41.666	1.00 14.76	AAAA
ATOM	561	CG	LEU	69	37.057				
ATOM	562	CDI	LEU	69	37.212	26.643	43.179	1.00 16.81	AAAA
			LEU	69	35.832	27.312	41.217	1.00 17.26	AAAA
ATOM .	563							1.00 15.11	AAAA
ATOM	564	С	LEU	69	39.623	27.796	38.969		
	565	0	LEU	69	39.500	28.934	38.504	1.00 13.30	AAAA
MOTA					40.803	27.204	39.090	1.00 13.40	AAAA
MOTA	566	N	MET	70					AAAA
ATOM	567	CA	MET	70	42.019	27.894	38.659	1.00 16.97	
	568	CB	MET	70	43.254	27.114	39.075	1.00 14.87	AAAA
ATOM .							40.582	1.00 15.18	AAAA
ATOM	569	CG	MET	70	43.335	26.886			
ATOM	570	SD	MET	70	44.828	25.954	41.060	1.00 28.71	AAAA
			MET	70	46.051	27.228	40.893	1.00 21.19	AAAA
ATOM	571	CE						1.00 19.11	AAAA
ATOM	572	С	MET	70	42.064	28.119	37.155		
	573	0	MET	70	42.498	29.170	36.700	1.00 17.10	AAAA
ATOM						27.118	36.389	1.00 15.06	AAAA
ATOM	574	N	GLI	71	41.648				AAAA
ATOM	575	CA	GLu	71	41.651	27.226	34.934	1.00 16.12	
			GLi	71	41.397	25.856	34.305	1.00 16.12	AAAA
MOTA	576	CB						1.00 20.26	AAAA
MOTA	577	CG	GLU	71	41.387	25.882	32.800		
	578	CD	GLU	71	42.782	25.920	32.193	1.00 32.31 [.]	AAAA
ATOM			•		42.893	25.741	30.958	1.00 27.07	AAAA
ATOM	579		GLU	71					
MOTA	580	OE2	GLU	71	43.762	26.117	32.941	1.00 24.85	AAAA
			GLU	71	40.580	28.208	34.466	1.00 16.48	AAAA
ATOM	581	С						1.00 17.20	· AAAA
ATOM .	582	0	GLU	71	40.831	29.066	33.611		
	583	N	ALA	72	39.380	28.097	35.027	1.00 15.68	AAAA
MOTA					38.300	28.998	34.644	1.00 16.07	AAAA
MOTA	584	CA	ALA	72					
ATOM	585	CB	ALA	72	37.035	28.669	35.425	1.00 17.21	AAAA
		c	ALA	72	38.678	30.453	34.897	1.00 19.07	AAAA
ATOM	586						34.054	1.00 15.92	AAAA
ATOM	587	0	ALA	72	38.448	31.326			
	588	N	GLU	73	39.260	30.726	36.062	1.00 15.86	AAAA
ATOM							36.372	1.00 15.50	AAAA
MOTA	589	CA	GLU	73	39.616	32.097			
	590	CB	GLU	73	40.046	32.210	37.828	1.00 14.12	AAAA
ATOM				73	40.430	33.615	38.214	1.00 14.24	AAAA
MOTA	591	CG	GLU					1.00 17.23	AAAA
ATOM ·	592	CD	GLU	73	40.961	33.699	39.629	1.00 17.43	
	593	OE1		73	40.147	33.696	40.573	1.00 18.51	AAAA
MOTE				73	42.201		39.793	1.00 20.88	AAAA
MOTA	594	OE2	∪نتی	13	42.201	33.753	2223		

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MOTA	59!	5 C	GLU	73	40.706	32.709	35.495	1.00 20.36	AAAA
ATOM	596		GLU		40.527	33.806	34.948	1.00 17.74	AAAA
ATOM	597				41.832	32.020		1.00 21.57	AAAA
MOTA	598				42.911			1.00 19.48	AAAA
ATOM	599				44.256			1.00 18.48	AAAA
MOTA	600				44.365			1.00 14.96	AAAA
MOTA	601				45.723			1.00 15.05	AAAA
MOTA	602				45.918			1.00 18.16	AAAA
MOTA	603				46.439			1.00 16.31	AAAA
MOTA MOTA	604 605		il ARG il ARG		46.843 46.466			1.00 19.74	AAAA
ATOM	606		ARG		42.643			1.00 14.73 1.00 16.86	AAAA
ATOM	. 607		ARG		43.148			1.00 15.41	
ATOM	608		CYS		41.859	_	32.547	1.00 17.56	AAAA AAAA
ATOM	609				41.544	31.833	31.115	1.00 18.24	AAAA
ATOM	610			75	41.474	30.414	30.545	1.00 20.94	AAAA
ATOM	611	. SG	CYS	75	43.047	29.514	30.572	1.00 19.30	AAAA
ATOM	612	С	CYS	75	40.216	32.561	30.898	1.00 15.81	AAAA
MOTA	613		CYS	75	39.762	32.748	29.762	1.00 17.79	AAAA
MOTA	614		GLN	76	39.601	32.959	32.007		AAAA
MOTA	615			76	38.339	33.686	32.010	1.00 23.22	AAAA
A'TOM	616			76 26	38.595	35.122	31.530	1.00 22.99	AAAA
ATOM	617 618			76 76	37.564 37.588	36.107	32.027	1.00 44.69	AAAA
MOTA MOTA	619		1 GLN	76	37.563	36.229 35.228	33.535 34.243	1.00 47.78	AAAA
ATOM	620		2 GLN	76	37.619	37.452	34.033	1.00 62.95 1.00 45.96	AAAA AAAA
ATOM	621	С	GLN	76	37.304	32.975	31.135	1.00 23.43	AAAA
ATOM	622	0	GLN	. 76	36.826	33.512	30.135	1.00 19.93	AAAA .
MOTA	623	N	CYS	7 7	36.951	31.754	31.521	1.00 15.97	AAAA
MOTA	624	CA		. 77	36.004	30.979	30.741	1.00 18.91	AAAA
ATOM	625	CB	CYS	77	36.738	30.225	29.623	1.00 24.64	AAAA
MOTA	626	SG	CYS	77	37.848	28.887	30.269	1.00 25.26	AAAA
MOTA	627 628	C	CYS CYS	77 77	35.302	29.951	31.594	1.00 19.68	AAAA
ATOM ATOM	629	N	VAL	78	35.685 34.254	29.702 29.366	32.732	1.00 20.02 1.00 16.00	AAAA
ATOM	630	CA	VAL	78	33.531	28.288	31.022 31.671	1.00 18.00	AAAA AAAA
ATOM	631	CB	VAL	78	32.016	28.455	31.557	1.00 15.57	AAAA
ATOM	632		l VAL	78	31.312	27.304	32.262	1.00 21.27	AAAA
MOTA	633	CG	Z VAL	78	31.603	29.792	32.151	1.00 19.47	AAAA
ATOM	634	С	VAL	78	33.950	27.077	30.859	1.00 24.02	AAAA
MOTA	635	0	VAL	78	33.499	26.894	29.718	1.00 24.08	AAAA
MOTA	636	N	PRO	79	34.848	26.249	31.420	1.00 18.91	AAAA
ATOM	637	CD	PRO	79	35.470	26.341	32.756	1.00 17.70	AAAA
ATOM	638	CA	PRO. PRO	79 79	35.320	25.056	30.720	1.00 23.37	AAAA
ATOM ATOM	639 640	CB	PRO	79 79	36.295 36.802	24.432 25.677	31.732 32.498	1.00 21.92 1.00 20.90	AAAA
ATOM	641	c	PRO	79	34.152	24.144	30.376	1.00 20.90	A AAA A AAA
ATOM	642	ō	PRO	79	33.177	24.064	31.119	1.00 22.20	AAAA
ATOM	643	N	LYS	80	34.245	23.488	29.224	1.00 23.35	AAAA
ATOM	644	CA	LYS	80	33.212	22.570	28.775	1.00 26.78	AAAA
MOTA	645	CB	LYS	80	33.708	21.853	27.518	1.00 32.33	AAAA
ATOM	646	CG	LYS	80	35.098	21.256	27.680	1.00 51.34	AAAA
ATOM	647	CD	LYS	80	35.669	20.817	26.336	1.00 68.70	AAAA
MOTA	648	CE.	LYS	80	37.131	20.401	26.451	1.00 70.04	AAAA
ATOM	649	NZ	LYS	80	37.688	19.949	25.141	1.00 73.72	AAAA
ATOM	650 65 <u>1</u>	C	LYS	80	32.875	21.571	29.875	1.00 24.71	AAAA
MOTA MOTA	652	Ŋ	LYS GLY	80 81	33.770 31.582	20.957 21.431	30.458 30.161	1.00 24.23 1.00 16.74	AAAA AAAA
ATOM	653	CA	GLY	81	31.126	20.509	31.194	1.00 18.74	AAAA
MOTA	654	C	GLY	81	31.151	21.039	32.630	1.00 18.38	AAAA
ATOM	655	ō	GLY	81	30.604	20.396	33.527	1.00 19.29	AAAA
ATOM	656	N	ALA	82	31.754	22.202	32.863	1.00 22.57	AAAA
MOTA	657	CA	ALA	82	31.858	22.738	34.235	1.00 20.65	AAAA
ATOM	658	CB	ALA	82	33.065	23.704	34.333	1.00 20.41	AAAA
ATOM	659	С	ALA	82	30.610	23.425	34.781	1.00 21.81	AAAA
MOTA	660	0	ALA	82	30.425	23.529	35.994	1.00 16.95	AAAA .

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- mov/	661	N	ARG	83	29.758	23.926	33.897	1.00 17.68	AAAA
MOTA									
MOTA	662	ÇA	ARG	83	28.549	24.596	34.360	1.00 15.04	
ATOM	663	CB	ARG	83	27.777	25.188	33.176	1.00 21.02	AAAA
MOTA	664	CG	ARG	83	26.938	26.395	33.528	1.00 36.77	AAAA
	665	CD	ARG	83	26.061	26.167	34.729	1.00 41.28	
ATOM					25.366	27.393	35.105	1.00 40.05	
ATCM	666	NE	ARG	83					
ATOM	667	CZ	_ARG	83	24.530	27.492	36.134	1.00 51.15	AAAA
MOTA	668	NHI	ARG	83	24.286	26.432	36.893	1.00 55.10	AAAA
ATCM	669	NHO	. ARG	83	23.931	28.646	36.399	1.00 54.26	AAAA
	670	С	ARG	83	27.701	23.530	35.030	1.00 21.33	
ATCM					27.193	23.708		1.00 24.88	
MOTA	671	0	ARG	83 -			36.130		
ATOM	672	N	GLU	84	27.565	22.406	34.352	1.00 18.76	•
ATOM	673	CA	GLU	84	26.768	21.299	34.859	1.00 24.12	AAAA
MOTA	674	CB	GLU	84	26.527	20.290	33.744	1.00 32.64	AAAA
ATOM	675	CG	GLU	84	27.769	19.994	32.925	1.00 37.91	AAAA
			GLU.	84	27.832	20.784	31.612	1.00 51.24	
MOTA	676	CD			-				
MOTA	677		. GLU	84	27.585	20.152	30.545	1.00 24.82	
MOTA	678	OE2	GLU	84	28.114	22.018	31.650	1.00 22.57	AAAA
ATOM	679	С	GLU	84	27.394	20.570	36.043	1.00 25.36	AAAA
MOTA	680	0	GLU	84	26.739	20.321	37.057	1.00 26.17	AAAA
	681	N	LYS	85	28.665	20.232	35.897	1.00 18.78	
MOTA					29.399	19.497	36.915	1.00 20.03	
MOTA	682	CA	LYS	85					
MOTA	683	CB	LYS	85	30.658	18.900	36.280	1.00 18.59	
MOTA	684	CG	LYS	85	31.603	18.223	37.268	1.00 35.69	
ATOM	685	CD	LYS	85	31.151	16.832	37.644	1.00 51.51	AAAA
ATOM	686	CE	LYS	85	31.451	15.864	36.520	1.00 59.18	AAAA
ATOM	687	NZ	LYS	85	32.914	15.858	36.240	1.00 56.63	AAAA
	688	c	LYS	85	29.811	20.263	38.181	1.00 18.31	AAAA
ATOM	689	o	LYS	85	29.696	19.738	39.290	1.00 21.65	
ATOM					30.274	21.495	38.012	1.00 19.45	
ATOM	690	N	TYR	86					
ATOM	691	CA	TYR ·		30.776	22.272	39.145	1.00 14.26	
ATOM	692	CB	TYR	86	32.207	22.692	38.840	1.00 14.95	
ATOM	693	CG	TYR	86	33.107	21.508	38.585	1.00 19.76	
ATOM	694	CD1	TYR	86 .	33.384	20.591	39.601	1.00 18.83	
ATOM	695	CE1	TYR	86	34.247	19.519	39.388	1.00 20.29	AAAA
ATOM	696	CD2		86	33.711	21.322	37.337	1.00 18.14	AAAA
MOTA	697	CE2		86	34.567	20.261	37.112	1.00 22.66	AAAA
	698	CZ	TYR	86	34.832	19.364	38.145	1.00 22.51	
ATOM					35.680	18.317	37.921	1.00 23.68	
atom	699	OH	TYR	86					
MOTA	700	С	TYR	86	29.967	23.493	39.526	1.00 19.03	
ATOM	701	0	TYR	86	30.353	24.226	40.450	1.00 19.18	
ATOM	702	N	ASN	87	28.873	23.721	38.803	1.00 17.59	
ATCM	703	CA	ASN	87	27.953	24.843	39.071	1.00 18.07	AAAA
ATCM	704	CB	ASN	87	27.413	24.730	40.514	1.00 23.87	AAAA
ATOM	705	CG	ASN	87	26.020	25.349	40.688	1.00 30.67	AAAA
	706		ASN	87	25.531	25.520	41.819	1.00 31.55	
ATCM				87	25.370	25.661	39.580	1.00 20.18	
ATOM	707		ASN					1.00 24.24	
ATOM	708	С	ASN	· 87	28.641	26.197	38.875		
ATOM	709	0	ASN	87	28.283	27.190	39.519	1.00 18.57	
ATCM	710	N	ILE	88	29.617	26.237	37.970	1.00 18.80	
ATCM	711	CA	ILE	88	30.353	27.471	37.680	1.00 18.55	AAAA
ATOM	712	CB	ILE	88	31.865	27.166	37.508	1.00 26.44	AAAA
	713		ILE	88	32.613	28.406	37.044	1.00 43.71	AAAA
ATCM		CG1		88	32.439	26.703	38.835	1.00 36.30	AAAA
	.714							1.00 24.08	AAAA
ATCM	715		ILE	88	32.295	27.735	39.888		
ATOM	716	С	ILE	88	29.887	28.142	36.392	1.00 14.36	
ATOM	717	0	ILE	88	29.584	27.459	35.426	1.00 21.93	AAAA
ATCM	718	N	GLY	89	29.843	29.473	36.380	1.00 18.71	AAAA
ATOM	719	CA	GLY	89	29.479	30.162	35.154	1.00 20.23	AAAA
	720	c	GLY	89	28.147	30.873	35.106	1.00 20.85	AAAA
ATOM	721	0	GLY	89	28.006	31.817	34.330	1.00 25.47	AAAA
ATCM				90	27.172	30.414	35.889	1.00 21.17	AAAA
ATCM	722	N	GLY					1.00 24.44	AAAA
ATOM	723	CA	GLY	90	25.863	31.060	35.898	1.00 20 50	
ATOM	724	C	GLY	90	25.862	32.371	36.668	1.00 30.60	
ATCM	725	0	GLY	90	26.900	32.788	37.168	1.00 28.13	AAAA
ATCM	726	N	TYR	91	24.708	33.036	36.755	1.00 23.38	AAAA
							-		•

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ATOM	727 CA TY	R 91	24.598	34.29	9 37.490	1.00 28.48	AAAA
MOTA	728 CB TY	₹ 91	23.144	34.75			
ATOM	729 CG TY	₹ 91	22.923				AAAA
ATOM	730 CD1 TY		23.329				AAAA
ATOM	731 CE1 TYP						AAAA
	732 CD2 TYF	01	23.130	38.250			AAAA
ATOM-			22.317				AAAA
ATOM	733 CE2 TYR		22.115				AAAA
MOTA	734 CZ TYR	91	22.521	38.002	40.327		AAAA
MOTA	735 OH TYR	91	22.306				
ATOM	736 C TYR	91	25.075		38.937		AAAA
ATOM	737 O TYR		25.713			1.00 23.59	AAAA
ATOM	738 N GLU		24.724				AAAA
ATOM	739 CA GLU						AAAA
			25.048				AAAA
ATOM			24.289	31.476	41.306		AAAA
ATOM	741 CG GLU		24.595	30.892	42.657		AAAA
ATOM	742 CD GLU		23.604	29.800	43.023	1.00 49.02	- AAAA
ATOM	743 OE1 GLU	92	24.008	28.829		1.00 45.51	
MOTA	744 OE2 GLU	92	22.418	29.931		1.00 38.16	AAAA
MOTA	745 C GLU	92	26.541	32.636		1.00 36.16	AAAA
ATOM	746 O GLU	92	27.045			1.00 25.78	AAAA
MOTA	747 N ASN	93		33.358		1.00 24.95	AAAA
ATOM	748 CA ASN	93	27.243	31.742		1.00 21.41	AAAA
			28.674	31.519		1.00 21.14	AAAA
ATOM	749 CB ASN	93	28.876	30.075	41.226	1.00 17.27	AAAA
MOTA	750 CG ASN	93	27.905	29.682	42.320	1.00 15.34	AAAA
ATOM	751 OD1 ASN	93	27.882	30.290	43.399	1.00 20.33	AAAA
ATOM	752 ND2 ASN	93	27.078	28.674	42.047	1.00 20.49	AAAA
MOTA	753 C ASN	93	29.378	31.778	39.445	1.00 22.25	
ATOM	754 O ASN	93	29.901	30.865	38.806	1.00 20.29	AAAA
ATOM	755 N PRO	94	29.451	33.057	39.045	1.00 25.45	AAAA
ATOM	756 CD PRO	94	29.027	34.221			AAAA
ATOM	757 CA PRO	94	30.055		39.839	1.00 23.03	AAAA
MOTA	758 CB PRO	94		33.523	37.794	1.00 23.05	AAAA
ATOM	759 CG PRO	94	29.669	35.004	37.759	1.00 28.71	AAAA
			28.528	35.112	. 38.755	1.00 40.02	AAAA
ATOM		94	31.554	33.384	37.697	1.00 26.51	AAAA
ATOM	761 O PRO	94	32.232	33.185	38.688	1.00 17.36	AAAA
ATOM	762 N VAL	95	32.068	33.498	36.478	1.00 21.12	AAAA
MOTA	763 CA VAL	95	33.506	33.493	36.281	1.00 17.00	AAAA
ATOM	764 CB VAL	95	33.851	33.242	34.796	1.00 25.15	AAAA
ATOM	765 CG1 VAL	95	35.326	33.537	34.533	1.00 27.19	
ATOM	766 CG2 VAL	95	33.551	31.791	34.443	1.00 27.19	AAAA
ATOM	767 C VAL	95	33.989	34.899			AAAA
ATOM	768 O VAL	95	33.426		36.686	1.00 17.42	AAAA
ATOM	769 N SER	96		35.894	36.237	1.00 23.43	AAAA
MOTA	770 CA SER	96	34.986	34.982	37.563	1.00 18.84	AAAA
			35.564	36.270	37.982	1.00 21.77	AAAA
MOTA		96	34.608	37.070	38.867	1.00 23.11	AAAA
ATOM	772 OG SER	96	34.723	36.679	40.223	1.00 24.43	AAAA
MOTA	773 C SER	96	36.835	35.987	38.789	1.00 29.09	AAAA
ATOM	774 O SER	96 .	37.117	34.828	39.115	1.00 27.12	AAAA
MOTA	775 N TYR	97	37.610	37.020	39.124	1.00 17.51	AAAA
MOTA	776 CA TYR	97	38.803	36.751	39.911	1.00 20.69	
ATOM	777 CB TYR	97	39.865	37.835	39.712		AAAA
ATOM	778 CG TYR	97		37.748		1.00 21.82	AAAA
ATOM	779 CD1 TYR	97			38.332	1.00 22.72	AAAA
ATOM	780 CEI TYR			38.414	37.235	1.00 28.47	AAAA
		97		38.265	35.949	1.00 24.45	AAAA
ATOM	781 CD2 TYR	97		36.929	38.112	1:00 19.74	AAAA
ATOM	782 CE2 TYR	97		36.771	36.832	1.00 21.63	AAAA
ATOM	783 CZ TYR	97	41.578	37.439	35.759	1.00 23.13	AAAA
ATOM	784 OH TYR	97	42.122	37.273	34.501	1.00 28.54	AAAA
ATOM	785 C TYR	97			41.393	1.00 20.12	AAAA
ATOM	786 O TYR	97		36.285		1.00 19.76	
MOTA	787 N ALA	98				1.00 19.76	AAAA
ATOM	788 CA ALA	98					AAAA
ATOM	789 CB ALA	98				1.00 22:23	AAAA
						1.00 27.92	AAAA
ATOM		98		34.743		1.00 23.56	AAAA
ATOM	791 O ALA	98				1.00 20.14	AAAA
MOTA	792 N MET	99	36.538	34.094	42.087	1.00 16.84	AAAA

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ATOM	793	CA	MEI	99	36.295	32.643	42.117	1.00 17.60	AAAA
ATOM	794	CB	MEI	99	35.864	32.137	40.736	1.00 17.05	AAAA
ATOM	795	CG	MET	99	36.999	31.824	39.793	1.00 11.16	AAAA
MOTA	796	SD	MET	99	36.314	31.698	38.113	1.00 16.54	AAAA
ATOM	797	CE	MET	99	35.165	30.295	38.312	1.00 17.83	AAAA
MOTA	798	С	MET	99	37.432	31.800	42.650	1.00 18.98	AAAA
ATOM	799	0	MET	99	37.197	30.753	43.251	1.00 18.21	AAAA
ATOM	. 800	N	PHE	100	38.670	32.216	42.420	1.00 12.87	AAAA
ATOM	801	CA	PHE	100	39.774	31.439	42.987	1.00 17.13	AAAA
ATOM	802	CB	PHE	100	40.559	30.681	41.917	1.00 15.23	AAAA
ATOM	803	CG	PHE	100	41.647	29.834	42.492	1.00 15.20	AAAA
ATOM	804	CD:	1 PHE	100	41.342	28.638	43.140	1.00 22.96	AAAA
ATOM	805	CD	2 PHE	100	42.972	30.282	42.488	1.00 17.12	AAAA
ATOM	806	CE	1 PHE	100	42:341	27.901	43.782	1.00 19.23	AAAA
ATOM	807	CE	2 PHE	100	43.974	29.552	43.129	1.00 16.99	AAAA
ATOM	808	CZ	PHE	100	43.658	28.360	43.779	1.00 17.78	AAAA
ATOM	809	С	PHE	100	40.755	32.305	43.774	1.00 20.54	AAAA
MOTA	810	0	PHE	100	41.088	31.990	44.912	1.00 21.45	AAAA
ATOM	811	N	THR	101	.41.219	33.401	43.187	1.00 18.02	AAAA
ATOM	812	CA	THR	101	42.177	34.245	43.902	1.00 15.25	AAAA
ATOM	813	CB	THR	101	42.715	35.341	42.976	1.00 16.33	AAAA
MOTA	814	OG:	THR	101	43.386	34.720	41.870	1.00 16.01	AAAA
ATOM	815	CG2	THR	101	43.706	36.226	43.697	1.00 16.31	AAAA
ATOM	816	С	THR	101	41.567	34.860	45.160	1.00 14.12	AAAA
MOTA	817	0	THR	101	42.110	34.707	46.244	1.00 16.86	AAAA
ATOM	818	И	GLY	102	40.435	35.541	45.008	1.00 13.77	AAAA
ATOM	819	CA	GLY	102	39.770	36.145	46.156	1.00 16.29	AAAA
ATOM	820	C	GLY	102	39.330	35.065	47.133	1.00 16.75	AAAA
MOTA	821	0	GĹY	102	39.502	35.202	48.338	1.00 14.48	AÄAA
MOTA	822	N	SER	103	38.752	33.986	46.615	1.00 16.24	AAAA
MOTA	823	CA	SER	103	38.315	32.890	47.488	1.00 16.72	AAAA
MOTA	824	CB	SER	103	37.567	31.821	46.684	1.00 15.97	AAAA
ATOM	825	OG	SER	103	36.339	32.349	46.197	1.00 26.86	AAAA
MOTA	826	С	SER	103	39.494	32.264	48.218	1.00 17.88	AAAA
MOTA	827	0	SER	103	39.405	31.974	49.419	1.00 14.17	AAAA
ATOM	828	N	SER	104	40.604	32.057	47.515	1.00 11.40	AAAA
ATOM	829	CA	SER	104	41.780	31.484	48.181	1.00 17.61	AAAA
ATOM	830	CB	SER	104	42.888	31.206	47.160	1.00 15.89	AAAA
ATOM	831	OG	SER	104	42.525	30.102	46.362	1.00 27.82	AAAA
ATOM	832	C	SER	104	42.332	32.404	49.271	1.00 17.02	AAAA
ATOM	833	O	SER	104 105	42.867 42.206	31.958	50.286	1.00 15.37	AAAA
ATOM	834	N CA	LEU	105	42.709	33.698	49.052	1.00 17.10 1.00 16.95	AAAA AAAA
ATOM	835 836	CB -		105	42.728	34.652 36.037	50.016 49.365	1.00 18.44	AAAA
ATOM	837	CG .	LEU	105	43.613	37.108	49.981	1.00 29.88	AAAA
ATOM	838		LEU	105	45.086	36 631	49.959	1.00 20.25	AAAA
ATOM ATOM	839		LEU	105	43.438	38 418	49.175	1.00 29.39	AAAA
ATOM	840	C	LEU	105	41.837	34 637	51.282	1.00 14.81	AAAA
ATOM	841	ō	LEU	105	42.334	34.703	52.404	1.00 17.74	AAAA
ATOM	842	N	ALA	106	40.532	34.531	51.095	1.00 19.28	AAAA
ATOM	843	CA	ALA	106	39.601	34.493	52.224	1.00 12.39	AAAA
ATOM	844	CB	ALA	106	38.140	34.574	51.704	1.00 11.58	AAAA
ATOM	845	c	ALA	106	39.807	33.210	53.023	1.00 14.79	AAAA
ATOM	846	ō	ALA	106	39.704	33.203	54.250	1.00 13.58	AAAA
ATOM	847	N	THR	107	40.114	32.128	52.318	1.00 13.67	AAAA
ATOM	848	CA	THR	107	40.314	30.819	52.956	1.00 13.21	AAAA
ATOM	849	CB	THR	107	40.187	29.708	51.902	1.00 14.95	AAAA
ATOM	850	OG1	THR	107	38.868	29.792	51.334	1.00 15.72	AAAA
ATOM	851	CG2	THR	107	40.422	28.311	52.511	1.00 9.51	AAAA
ATOM	852	C	THR	107	41.649	30.751	53.687	1.00 15.80	AAAA
ATOM	853	ō	THR	107	41.734	30.206	54.792	1.00 15.63	AAAA
ATOM	854		GLY	108	42.696	31.294	53.082	1.00 14.08	AAAA
ATOM	855	CA	GLY	108	43.968	31.298	53.765	1.00 14.62	AAAA
ATOM ·	856	Ç.	GLY	108	43.801.	32.119	55.041	1.00 20.05	AAAA
ATOM	857	ö	GLY	108	44.417	31.813	56.063	1.00 17.53	AAAA
ATOM	858	N	SER	109	42.963	33.158	54.988	1.00 15.26	AAAA
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MOTA	859 CA	SER 109	42.72	7 34.020	56.153	1.00 14.54	
MOTA	860, CB	SER 109	41.906				AAAA
ATOM	861 OG	SER 109				1.00 15.58	AAAA
ATOM	862 C	SER 109	42.037		57.297		. AAAA
MOTA	863 O	SER 109	42.189		58.487		AAAA
ATOM	864 N	THR 110	41.261	32.247			AAAA
MOTA	865 CA	THR 110	40.608				AAAA
ATOM	866 CB	THR 110	39.452				AAAA
ATOM	867 OG1		38.346				AAAA
ATOM .		THR 110	39.061			_	AAAA
MOTA	869 C	THR 110	41.633		•	1.00 12.91	AAAA
MOTA		THR 110	41.574			1.00 18.44	AAAA
ATOM		VAL 111	42.584		59.806	1.00 16.30	AAAA
ATOM	_	VAL 111	43.614		57.816	1.00 15.20	AAAA
ATOM		VAL 111	44.517			1.00 20.45	AAAA
ATOM	874 CG1			28.514	57.323	1.00 20.02	AAAA
ATOM	875 CG2		45.652	27.765	58.005	1.00 21.79	AAAA
ATOM		VAL 111	43.697		56.482	1.00 19.07	AAAA
ATOM			44.456	30.075	59.327	1.00 18.21	AAAA
. ATOM			44.838	29.672	60.431	1.00 18.65	AAAA
ATOM			44.731	31.302	58.890	1.00 16.82	AAAA
		GLN 112	45.493	32.232	59.719	1.00 20.13	AAAA
MOTA		GLN 112	45.751	33.540	58.970	1.00 22.39	AAAA
ATOM		GLN 112	46.593	33.360	57.723	1.00 21.17	AAAA
ATOM		GLN 112	46.797	34.651	56.982	1.00 24.82	AAAA
ATOM	883 OE1 C		47.772	35.381	57.219	1.00 25.62	AAAA
ATOM		SLN 112	45.866	34.963	56.091	1.00 13.16	AAAA
ATOM		SLN 112	44.743	32.516	61.012	1.00 23.99	AAAA
MOTA		3LN 112	45.340	32.593	62.079	1.00 17.94	AAAA
ATOM		ALA 113	43.431	32.700	60.924	1.00 15.60	AAAA
ATOM		LA 113	42.653		62.138	1.00 15.04	AAAA
ATOM		LA 113	41.191	33.138	61.802	1.00 18.65	AAAA
ATOM		LA 113	42.807	31.751	63.083	1.00 14.84	AAAA
ATOM		LA 113	42.941	31.909	64.296	1.00 21.05	AAAA
MOTA		LE 114	42.767	30.550	62.534	1.00 16.45	AAAA ·
ATOM		LE 114	42.919	29.383	63.389	1.00 15.38	AAAA
ATOM		LE 114	42.600	28.100	62.637	1.00 15.22	AAAA
ATOM	895 CG2 I		42.888	26.893	63.537	1.00 15.72	AAAA
ATOM	896 CG1 II		41.110	28.112	62.244	1.00 19.28	AAAA
ATOM	897 CD1 II		40.744	27.038	61.191	1.00 13.43	AAAA
ATOM ATOM		LE 114	44.329	29.318	63.968	1.00 18.02	AAAA
ATOM		LE 114	44.508	28.998	65.156	1.00 20.38	AAAA
		LU 115	45.328	29.629	63.144	1.00 15.27	AAAA
ATOM	901 CA GI		46.726	29.625	63.614	1.00 21.48	AAAA
ATOM	902 CB GI		47.690	30.080	62.506	1.00 21.76	AAAA
MOTA	903 CG GI		47.884	29.080	61.386	1.00 15.78	AAAA
MO. A	904 CD GI		.48.670	29.648	60.211	1.00 20.04	AAAA
MO7 K	905 OE1 GL		49.051	30.843	60.239	1.00 21.48	AAAA
A::OM	906 OE2 GL		48.901	28.902	59.241	1.00 26.59	AAAA
ATOM	907 C GL		46.877	30.559		1.00 23.55	AAAA
ATOM	908 O GL		47.509	30.212	65.815	1.00 23.03	AAAA
ATOM	909 N GL				64.703	1.00 22.73	AAAA
ATOM	910 CA GL			32.735	65.774	1.00 20.54	AAAA
ATOM	911 CB GL			34.044	65.320	1.00 18.40	AAAA
MOTA	.912 CG GL		46.562			1.00 19.76	AAAA
ATOM	913 CD GL					1.00 27.24	AAAA
ATOM	914 OE1 GL		48.164			1.00 18.44	AAAA
ATOM	915 CE2 GL		48.919	34.543		1.00 23.17	AAAA
ATOM	916 C GLU					1.00 25.39	AAAA
atom	917 0 GLI			~		1.00 22.87	AAAA
MOTA	918 N PHE	E 117				.00 18.78	AAAA
MOTA	919 CA PHE	117				.00 22.11	•
MOTA	920 CB PHE	117				00 22.11	AAAA
MOTA	921 CG PHE	117				.00 24.06	AAAA
MOTA	922 CD1 PHE	117					AAAA
MOTA	923 CD2 PHE					.00 19.67 .00 24.08	AAAA
MOTA	924 CE1 PHE						AAAA
		·			A. 190 T	.00 23.68	AAAA

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PCT/US00/24700

ATOM		CE2 PH	E 117	40.79	9 28.26	2 70.156	1.00 24.04	Aaaa
MOTA	926	CZ PH	117	40.07	8 29.17			
ATOM	927	C PHE	117	44.58				AAAA
ATOM		O PHE		44.61				AAAA
MOTA		_						AAAA
			_	45.23				AAAA
MOTA		CA LEU		46.02	5 28.113	3 68.549	1.00 20.73	AAAA
MOTA	931-	CB LEU	118	46.358	3 27.07	5 67.480	1.00 17.90	AAAA
ATOM	932	CG LEU	118	45.148	3 26.264			
ATOM	933	CD1 LEU		45.591				AAAA
ATOM		CD2 LEU		44.520			1.00 34.23	AAAA
ATOM								AAAA
		C LEU		47.290			1.00 26.49	AAAA
ATOM) LEU		47.908			1.00 26.34	AAAA
MOTA	937 1	1 LYS	119	47.672	29.848	8 68.975		AAAA
ATOM	938 (CA LYS	119	48.835			1.00 28.53	
ATOM	939 (B LYS		49.392			1.00 20.55	AAAA
ATOM		G LYS					1.00 30.15	AAAA
			-	49.915				· AAAA
ATOM		D LYS		50.291			1.00 28.98	AAAA
ATOM		E LYS	119	50.905		65.380	1.00 31.07	AAAA
ATOM	943 N	IZ LYS	119	51.195	33.551	64.745	1.00 22.46	AAAA
ATOM	944 C	LYS	119	48.335			1.00 35.74	
ATOM	945 C		119	49.117			1.00 33.74	AAAA
MOTA	946 N		120				1.00 27.10	AAAA
				47.018			1.00 25.20	AAAA
ATOM		A GLY	120	46.445			1.00 30.18	AAAA
ATOM	948 C		120	45.913	33.007	72.122	1.00 31.91	AAAA
MOTA	949 O	GLY	120	45.540	33.665	73.094	1.00 34.76	AAAA
ATOM	950 N	ASN	121	45.889	33.495		1.00 20.56	AAAA
ATOM	951 C	A ASN	121	45.353	34.825			
ATOM	952 C		121	46.278	35.634		1.00 25.58	AAAA
ATOM	953 C		121				1.00 29.99	AAAA
ATOM -				47.641	35.827		1.00 24.43	AAAA
		D1 ASN	121	48.396	34.874	70.588	1.00 54.63	AAAA
ATOM		D2 ASN	121	47.944	37.045	70.817	1.00 41.69	AAAA
MOTA	956 C	ASN	121	43.941	34.759	70.135	1.00 18.85	AAAA
MOTA	957 O	ASN	121	43.421	33.675	69.899	1.00 24.77	AAAA
ATOM	958 N.	VAL	122	43.310	35.918	69.991		
ATOM	959 C		122	41.936	35.994	69.499	1.00 19.55	AAAA
ATOM	960 CI						1.00 22.90	AAAA
			122	41.053	36.832	70.449	1.00 31.47	AAAA
ATOM		1 VAL	122	39.649	37.006	69.851	1.00 31.52	AAAA
MOTA		2 VAL	122	40.986	36.154	71.810	1.00 32.50	AAAA
ATOM	963 C	VAL	122	41.953	36.632	68.130	1.00 16.87	AAAA
ATOM	964 O	∇AL	122	42.518	37.710	67.938	1.00 24.08	AAAA
ATOM	965 N	ALA	123	41.321	35.983	67.159		
ATOM	966 CA		123	41.360			1.00 18.67	AAAA
ATOM	967 CE				36.532	65.821	1.00 10.18	AAAA
			123	42.346	35.743	64.990	1.00 19.04	AAAA
ATOM	968 C	ALA	123	40.000	36.551	65.131	1.00 13.72	AAAA
MOTA	969 O	ALA	123	39.108	35.761	65.439	1.00 20.78	AAAA
ATOM	970 N	PHE	124	39.871	37.457	64.180	1.00 12.92	AAAA
ATOM	971 CA	PHE	124	38.649	37.610	63.401	1.00 14.67	
ATOM	972 CB	PHE	124	37.904	38.878	63.85€		AAAA
MOTA	973 CG		124	36.660	39.209		1.00 14.67	AAAA
ATOM		1 PHE				63.049	1.00 20.28	AAAA
			124	35.811	38.209	62.587	1.00 18.56	AAAA
ATOM		2 PHE	124	36.286	40.545	62.843	1.00 19.53	AAAA
MOTA		1 PHE	124	34.609	38.532	61.937	1.00 18.75	AAAA
ATOM	977 CE	2 PHE	124	35.072	40.875	62.193	1.00 20.18	AAAA
ATOM	978 CZ	PHE	124	34.242		61.744	1.00 21.57	AAAA
ATOM	979 C	PHE	124	39.016	37.712			
	980 0					61.930	1.00 22.60	AAAA
MOTA		PHE	124	39.823	38.558	61.535	1.00 19.22	AAAA
ATOM	981 N	ASN	125	38.449	36.820	61.126	1.00 19.39	AAAA
ATOM	982 CA	ASN	125	38.651	36.858	59.691	1.00 16.80	AAAA
ATOM	983 CB	ASN	125	39.122	35.507	59.150	1.00 15.71	AAAA
ATCM	984 CG	ASN	125	39.063	35.469	57.649	1.00 12.84	
ATOM		ASN	125	39.216	36.508			AAAA
						57.006	1.00 14.91	AAAA
ATOM		ASN	125	38.853	34.272	57.065	1.00 16.21	AAAA
ATCM	987 C	ASN	125	37.315	37.210	59.038	1.00 16.22	AAAA
ATOM	988 0	ASN	125	36.502	36.330	58.755	1.00 15.28	AAAA
ATOM	989 N	PRO	126	37.071	38.502	58.775	1.00 14.84	AAAA
ATOM	990 CD	PRO	126	37.908	39.684	59.052	1.00 18.10	AAAA
				5500	004	22.032	T.00 TO.10	AAAA
								•

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ATOM			CA PR		35	. 811	38.910	58.15	5 1.0	0 17.33	АААА
ATOM			B PR		35	.912	40.434			0 16.32	AAAA
ATOM			G PRO		37	416	40.655	5 58.008		0 20.95	AAAA
ATOM						549	38.359	56.752		0 13,78	AAAA
ATOM						404	38.291			0 17.03	AAAA
ATOM						607	37.972			0 14.57	AAAA
ATOM			A AL			463	37.443		1.0	0 17.37	AAAA
ATOM			B ALA			816	37.540			0 14.48	AAAA
ATOM						982	35.998		1.0	0 19.77	AAAA
MOTA MOTA						490	35.500			0 15.62	· AAAA
ATOM				-		111	35.339			0 13.54	AAAA
ATOM					_	725	33.939			0 13.53	AAAA
ATOM						234 414	33.679			0 14.49	AAAA
ATOM							34.585			0 15.65	AAAA
ATOM						883 487	32.420		1.00	0 13.35	AAAA
ATOM	100		GLY		31.		32.058			16.28	AAAA -
ATOM	100	_	GLY		. 30.		31.831			15.69	AAAA
ATOM	100		MET		32.		32.021 31.448			16.10	AAAA
ATOM	101				31.		31.163			15.00	AAAA
ATOM	101			130	32.		31.215			13.35	AAAA
ATOM	1013	2 CG		130	33.		32.573	51.731		17.03	AAAA
MOTA	1013	3 SI	MET	130	34.		32.877	50.425		15.41	AAAA
MOTA	1014	4 CE	MET	130	33.		32.973	49.073		46.82	AAAA AAAA
MOTA	1015		MET	130	31.3	296	29.756	52.885		12.49	AAAA
ATOM	1016		MET	130	31.		28.789	52.297		19.54	AAAA
ATOM	1017		HIS	131	30.:		29.695	53.617		16.24	AAAA
MOTA	1018			131	29.5		28.448	54.014		13.80	AAAA
ATOM	1019			131	28.7		28.694	55.316		15.91	AAAA
ATOM ATOM	1020 1021		HIS 2 HIS	131	. 27.6		29.625	55.175		13.08	AAAA
ATOM	1021		1 HIS	131 131	26.7		30.063	56.096		12.46	AAAA
ATOM	1023		1 HIS	131	27.2		30.190	53.976		22.48	AAAA
ATOM	1024		2 HIS	131 .	26.1 25.8		30.936 30.875	54.166		16.56	AAAA
ATOM	1025		HIS	131	28.6		27.663	55.442 53.066		23.56	AAAA
ATOM	1026		HIS	131	28.1		26.658	53.470		13.69 17.21	AAAA
MOTA	1027	N	HIS	132	28.5		28.115	51.830		14.51	AAAA
MOTA	1028	CA	HIS	132	27.6		27.400	50.887		20.19	AAAA AAAA
ATOM	1029	CB	HIS	132	26.8		28.416	50.054		17.26	AAAA
ATOM	1030	CG	HIS	132	25.7		29.070	50.810		16.85	AAAA
MOTA	1031		? HIS	132	24.7	87 2	28.542	51.604		13.74	AAAA
ATOM	1032		HIS	132	25.4		30.424	50.756		24.80	AAAA
ATOM	1033		HIS	132	24.4		30.700	51.486	1.00	12.68	AAAA
ATOM	1034		HIS	132	23.9		29.576	52.010	1.00	28.65	AAAA
ATOM ATOM	1035 1036	C	HIS HIS	132	28.3		6.412	49.946		16.89	AAAA
ATOM	1037	.:J	ALA	132 133	27.7		5.487	49.460		14.58	AAAA
ATOM	1038	.v CA	ALA	133	29.66 30.3		6.580	49.689		16.79	AAAA
ATOM	1039	CB	ALA	133	31.7		5.680 6.194	48.740		13.76	AAAA
ATOM	1040	c	ALA	133	30.4		4.219	48.412 49.179		14.95	AAAA
ATOM	1041	ō	ALA	133 .	30.55		3.939	50.355		18.80 16.86	AAAA
ATOM	1042	:1	PHE	134	30.30		3.306	48.209		13.76	AAAA
MOTA	1043	CA	PHE	134	30.37		1.868	48.451		19.77	AAAA · AAAA
ATOM	1044	CB	PHE	134	29.31		1.132	47.620		15.59	AAAA
ATOM	1045	CG	PHE	134	27.91		1.525	47.975		17.22	AAAA
MOTA	1046	CD1	PHE	134	27.13		2.259	47.091		17.88	AAAA
MOTA	1047		PHE	134	27.39	2 2	1.187	49.222		21.68	AAAA
MOTA	1048	CE1		134	25.83		2.653	47.445		23.07	AAAA
MOTA	1049	CE2	PHE	134	26.09		1.578	49.585		17.64	AAAA
MOTA	1050	cz	PHE	134	25.32		2.308	48.696		19.71	AAAA
MOTA	1051	5	PHE	134	31.76	_	1.354	48.098		14.76	AAAA
MOTA	1052	5	PHE	134	32.54		2.049	47.442	1.00	18.05	AAAA
MOTA	1053		LYS	135	32.06	_	0.124	48.515	1.00		AAAA
ATOM	1054		LYS	135	33.36		9.551	48.269	1.00		AAAA
MOTA	1055		LYS	135	33.36		3.070	48.699	1.00		AAAA
MOTA	1056	CG .	LYS	135	34.64	U 17	7.300	48.400	1.00	30.43	AAAA

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Figure 16-17

		_							
ATOM	1057	CD	LYS	135	34.597	15.867	48.977	1.00 30.26	AAAA
ATOM	1058	CE	LYS	135	34.862	15.805	50.486	1.00 35.01	AAAA
ATOM	1059		LYS	135	36.304	16.023	50.895	1.00 20.61	AAAA
ATOM	1060		LYS	135	33.854	19.687	46.836	1.00 16.60	AAAA
ATOM	1061	. 0	LYS	135	35.020	20.020	46.584	1.00 17.24	AAAA
ATOM	1062	N	SER	136	32.944	19.483	45.893	1.00 18.01	AAAA
MOTA	1063		SER	136	33.301	19.528	44.490	1.00 15.26	
									AAAA
MOTA	1064		SER	136	33.339	18.094	43.940	1.00 18.07	AAAA
ATOM	1065	OG	SER	136	34.135	17.261	44.762	1.00 22.22	AAAA
ATOM	1066	С	SER	136	32.345	20.355	43.658	1.00 15.40	AAAA
MOTA	1067		SER	136	32.162	20.071	42.475	1.00 18.77	AAAA
ATOM	1068	N	ARG	137	31.754	21.401	44.237	$\cdot 1.0019.71$	AAAA
· ATOM	1069	CA	ARG	137	30.805	22.216	43.482	1.00 17.29	AAAA
ATOM	1070	CB	ARG	137	29.481	21.448	43.366	1.00 24.19	AAAA
MOTA	1071		ARG	137	28.290	22.273	42.937	1.00 32.56	AAAA
ATOM	1072	CD	ARG	137	27.026	21.424	42.980	1.00 47.98	AAAA
MOTA	1073	NE	ARG	137	26.951	20.493	41.862	1.00 50.95	AAAA
ATOM	1074		ARG	137	26.392	20.781	40.691	1.00 50.38	AAAA
ATOM	1075	NH1		137	.25.854	21.976	40.485	1.00 45.26	AAAA
MOTA	1076	NH2	ARG	137	26.375	19.876	39.722	1.00 55.31	AAAA
MOTA	1077	C	ARG	137	30.537	23.595	44.095	1.00 16.14	AAAA
ATOM	1078		ARG	137	30.439	23.711	45.308	1.00 16.88	
									AAAA
MOTA	1079		ALA	138	30.395	24.621	43.252	1.00 18.07	AAAA
MOTA	1080	CA.	ALA	138	30.117	25.976	43.735	1.00 21.48	AAAA
MOTA	1081	CB .	ALA	138	30.460	27.024	42.631	1.00 16.55	AAAA
ATOM	1082		ALA	138	28.642	26.090	44.135	1.00 21.04	AAAA
					27.798				
MOTA	1083		ALA	138		25.339	43.641	1.00 18.97	AAAA
ATOM	1084	N .	ASN	139	28.321	27.019	45.029	1.00 13.83	AAAA
ATOM	1085	CA Z	ASN	139	26.952	27.158	45.468	1.00 12.92	AAAA
ATOM	1086		ASN	139	26.566	25.899	46.274	1.00 13.14	AAAA
			ASN	139	25.162				
MOTA	1087					25.961	46.832	1.00 20.34	AAAA
ATOM	1088	OD1	ASN	139	24.186	26.068	46.086	1.00 19.76	AAAA
ATOM	1089	ND2	ASN	139	25.048	25.881	. 48.157	1.00 16.36	AAAA
ATOM	1090	C i	ASN	139	26.756	28.409	46.315	1.00 20.92	AAAA
	1091			139	27.603	28.738			
MOTA			ASN				47.148	1.00 16.81	AAAA
ATOM	1092		GLY	140	25.644	29.105	46.086	1.00 19.30	AAAA
MOTA	1093	CA (3LY	140	25.330	30.295	46.864	1.00 21.34	AAAA
ATOM	1094	C C	GLY	140	26.393	31.378	46.888	1.00 20.19	AAAA
ATOM	1095		SLY	140	26.653	31.968	47.943	1.00 18.77	AAAA
ATOM	1096		PHE	141	26.996	31.649	45.733	1.00 15.52	AAAA
MOTA	1097	CA I	PHE	141	28.034	32.675	45.600	1.00 20.71	AAAA
ATOM	1098	CB E	PHE	141	27.711	33.952	46.388	1.00 20.03	AAAA
ATOM	1099	CG I	PHE	141	26.355	34.544	46.127	1.00 28.32	AAAA
	1100	CD1 F		141	25.855	35.526	46.997	1.00 24.25	AAAA
ATOM									
ATOM	1101	CD2 F		141	25.589	34.170	45.029	1.00 30.11	AAAA
ATOM	1102	CE1 F	PHE	141	24.628	36.116	46.775	1.00 25.94	AAAA
ATOM	1103	CE2 P	HE	141	24.346	34.766	44.801	1.00 21.€:	AAAA
ATOM	1104		HE	141	23.870	35.741	45.677	1.00 24.4	AAAA
ATOM	1105		HE	141	29.357	32.188	46.158	1.00 14.45	AAAA
ATOM	1106	O P	HE	141	30.336	32.914	46.111	1.00 16.39	AAAA
ATOM	1107	N C	YS	142	29.389	30.982	46.716	1.00 16.77	AAAA
	1108	CA C		142	30.629	30.466	47.285	1.00 17.71	AAAA
ATOM				_	30.347				
ATOM	1109		YS	142		29.845	48.659	1.00 13.95	AAAA
MOTA	1110	SG C	YS	142	29.606	30.985	49.846	1.00 16.63	AAAA
MOTA	1111	C C	YS	142	31.313	29.421	46.401	1.00 18.09	AAAA
	1112		YS	142	30.647	28.527	45.856	1.00 16.60	AAAA
MOTA									
MOTA	1113		YR	143	32.639	29.539	46.272	1.00 12.50	AAAA
ATOM	1114	CA T	YR	143	33.429	28.603	45.478	1.00 15.32	AAAA
ATOM	1115	CB T	YR	143	34.333	29.322	44.473	1.00 13.07	AAAA
	1116		YR	143	33.614	30.338	43.612	1.00 15.80	AAAA
MOTA									
ATOM	1117	CD1 T		143	33.396	31.636	44.071	1.00 15.48	AAAA
MOTA	1118	CE1 T	YR	143	32.740	32.589	43.270	1.00 11.99	AAAA
ATOM	1119	CD2 T	YR	143	33.157	29.999	42.336	1.00 14.60	AAAA
		CE2 T		143	32.501	30.935	41.532	1.00 10.74	AAAA
MOTA	1120								
ATOM	1121		YR	143	32.301	32.229	42.008	1.00 20.89	AAAA
ATOM	1122	OH T	YR	143	31.698	33.177	41.208	1.00 18.87	AAAA
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ATOM			TY	143	34.31	0 27.72	3 46.358	3 1.00 17.35	
ATOM			TYF	₹ 143	34.58			1.00 16.67	AAAA
ATOM	1 112	25 N	ILE	144	34.76				
ATOM	112	26 C	A ILE	144	35.59				AAAA
ATOM	112	7 C	B ILE		37.01				AAAA
ATOM			G2 ILE		37.86				AAAA
ATOM			G1 ILE			_			AAAA
MOTA			D1 ILE		37.61				AAAA
		-		_	39.05			1.00 17.42	AAAA
ATOM					34.95			1.00 17.22	AAAA
ATOM		_			34.60	6 28.71	6 50.220	1.00 14.72	AAAA
ATOM					34.79	8 26.48	6 50.474		AAAA
MOTA			a asn	145	34.17	0 26.49			
ATOM			B ASN	145	33.40				AAAA
ATOM	113	6 C	g asn	145	32.42				AAAA
MOTA	113	7 0	D1 ASN	145	32.80				AAAA
ATOM	113	B NI	D2 ASN	145	31.17				AAAA
MOTA	113:		ASN	145	35.26				AAAA
MOTA	114		ASN	145	35.812				AAAA
ATOM	114		ASN	146	35.599				AAAA
ATOM	1142			146				1.00 12.34	AAAA
ATÓM	1143				36.685				AAAA
ATOM	1144			146	37.161	29.464		1.00 15.81	AAAA
				146	36.101			1.00 15.25	AAAA
ATOM	1145		1 ASN	146			56.034	1.00 13.57	AAAA
MOTA	1146		2 ASN	146	35.156		53.996	1.00 10.85	AAAA
ATOM	1147		ASN	146	36.306		55,613	1.00 13.04	AAAA
ATOM	1148		ASN	146	37.160	26.865	56.314	1.00 14.76	AAAA
ATOM	1149		PRO	147	35.025	27.489		1.00 14.28	AAAA
ATOM	1150		PRO	147	33.817	28.175		1.00 7.62	AAAA
ATOM	1151	CA	PRO	147	34.750			1.00 13.51	
MOTA	1152	CB	PRO	147	33.251			1.00 14.44	AAAA
MOTA	1153	CG	PRO	147	33.056			1.00 12.32	AAAA
MOTA	1154	С	PRO	147	35.118			1.00 18.86	AAAA
ATOM	1155	0	PRO	147	35.678		58.251		AAAA
ATOM	1156	N	ALA	148	34.818			1.00 16.24	AAAA
ATOM	1157	CA	ALA	148	35.122	23.200		1.00 15.01	AAAA
ATOM	1158	CB	ALA	148	34.402	22.561	56.080	1.00 15.58	AAAA
ATOM	1159	C	ALA	148	36.624		54.882	1.00 12.93	AAAA
ATOM	1160	ō	ALA	148	37.138	22.956	55.984	1.00 14.94	AAAA
ATOM	1161	N	VAL	149		21.999	56.560	1.00 14.69	AAAA
ATOM	1162	CA	VAL	149	37.328	23.817	55.263	1.00 12.49	AAAA
ATOM	1163	СВ	VAL	149	38.778	23.708	55.163	1.00 15.31	AAAA
ATOM	1164		VAL		39.364	24.797	54.243	1.00 14.77	AAAA
MOTA	1165		VAL	149	40.899	24.870	54.369	1.00 14.68	AAAA
ATOM	1166	C		149	38.981	24.501	52.808	1.00 12.50	AAAA
			VAL	149	39.323	23.887	56.572	1.00 20.14	. AAAA
ATOM	1167	0	VAL	149	40.172	23.109	57.028	1.00 17.32	AAAA
ATOM	1168	N	GLY	150	38.815	24.899	57.271	1.00 15.45	AAAA
ATOM	1169	CA	GLY	50	39.284	25.168	58.622	1.00 20.96	AAAA
ATOM	1170	C	GLY	- 50	39.030	24.053	59.621	1.00 24.16	AAAA
MOTA	1171	0	GLY	150	39.888	23.738	60.458	1.00 19.50	AAAA
ATOM	1172	N	ILE	151	37.842	23.465	59.557	1.00 16.67	AAAA
ATOM	1173	CA	ILE	151	37.490		60.461	1.00 19.56	
MOTA	1174	CB	ILE	151	35.992	22.052	60.348	1.00 16.46	AAAA
ATOM	1175	CG2	ILE	151	35.667	20.709	61.036	1.00 17.93	AAAA
ATOM	1176	CG1	ILE	151	35.180	23.209	60.959	1.00 17.33	AAAA
MOTA	1177		ILE	151	33.686	23.123			AAAA
MOTA	1178	C	ILE	151 .	38.352	21.148	60.672	1.00 18.71	AAAA
ATOM	1179	Ō	ILE	151	38.796	20 422	60.164	1.00 22.66	AAAA
ATOM	1180	N	GLU	152		20.472	61.087	1.00 20.08	aaaa
	1181	CA			38.599	20.861	58.888	1.00 19.71	AAAA
	1182		GLU	152	39.434	19.718	58.533	1.00 13.85	AAAA
ATOM	1182	CB	GLU	152	39.362	19.437	57.033	1.00 20.21	AAAA
ATOM		CG .	GLU	152	38.033	18.833	56.624	1.00 22.16	AAAA
ATOM	1184	CD	GLU	152	37.838	17.430	57.166	1.00 26.94	AAAA
ATOM	1185	0E1		152	36.720	16.906	57.035	1.00 25.03	AAAA
ATCM .	1186	CE2		152	38.800	16.846		1.00 24.95	AAAA
ATOM	1187		GLU	152	40.865	20.010		1.00 16.85	AAAA
atom	1188	0	GLU '	152	41.629	19.110		1.00 19.25	AAAA
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						• .					
MOTA	1189	N	.TYR	153		41.228	21.290	58.931	1.00	14.74	AAAA
ATOM	1190	CA	TYR	153		42.574	21.672	59.350		17.71	AAAA
ATOM	1191	СВ	_			42.757	23.193	59.179		13.26	AAAA
ATOM	1192	CG				44.059	23.727	59.729		16.36	AAAA
MOTA	1193		1 TYR			45.234	23.726	58.967		18.41	
			1 TYR								AAAA
ATOM	1194					46.438	24.219	59.511		21.03	AAAA
ATOM	1195		2 TYR			44.115	24.220	61.028		21.16	AAAA
MOTA	1196		2 TYR			45.288	24.705	61.570		19.76	AAAA
ATOM	1197	CZ	TYR			46.440	24.711	60.824	1.00	25.97	AAAA
ATOM	1198	ОН	TYR			47.571	25.235	61.410	1.00	23.15	AAAA
ATOM	1199	C	TYR	153	-	42.712	21.274	60.828	1.00	20.00	AAAA
ATOM	1200	0	TYR	153		43.722	20.698	61.247	1.00	19.61	AAAA
ATOM	1201	N	LEU	154		41.683	21.569	61.616	1.00	17.78	ÄAAA
MOTA	1202	CA	LEU	154		41.698	21.239	63.042	1.00	17.26	AAAA
MOTA	1203	CB	LEU	154		40.511	21.913	63.744		20.44	AAAA
ATOM	1204	CG	LEU			40.636	23.434	63.942		19.57	AAAA
ATOM	1205		LEU	154		39.277	24.046	64.309		22.48	AAAA
MOTA	1206		2 LEU	154		41.692	23.709	65.044		20.84	AAAA
MOTA	1207	C C	LEU	154		41.669	19.715	63.262		19.69	
	1208	ō	LEU	154		42.357	19.191	64.149			AAAA
ATOM		Ŋ		155						22.91	AAAA
ATOM	1209		ARG			40.878	18.996	62.469		20.88	-
MOTA	1210	CA	ARG	155		40.840	17.539	62.622		22.64	AAAA
MOTA	1211	CB	ARG	155		39.829	16.905	61.652		25.69	AAAA
MOTA	1212	CG	ARG	155		38.384	17.394	61.893		27.64	AAAA
MOTA	1213	CD	ARG	155		37.382	16.834	60.892		25.67	AAAA
MOTA	1214	NE	ARG	155		36.931	15.497	61.246		30.88	AAAA
MOTA	1215	CZ	ARG	155		36.135	14.753	60.488	1.00	36.28	AAAA
ATOM	1216	NH1	ARG	155	•	35.705	15.218	59.318	1.00	26.96	AAAA
MOTA	1217	NH2	ARG	155		35.737	13.562	60.923	1.00	27.33	AAAA
ATOM	1218	С	ARG	155		42.235	16.966	62.390	1.00	28.00	AAAA
MOTA	1219	0	ARG ·	155		42.674	16.070	63.119	1.00	28.05	AAAA
MOTA	1220	N	LYS	156		42.949	17.486	61.395	1.00	23.53	AAAA
MOTA	1221	CA	LYS	156		44.290	16.977	61.128	1.00	26.79	AAAA
MOTA	1222	ÇВ	LYS	156		44.854	17.558	59.824	1.00	26.01	AAAA
ATOM	1223	CG	LYS	156		46.213	16.955	59.444	1.00	29.70	AAAA
ATOM	1224	CD	LYS	156		46.632	17.308	58.035		28.77	AAAA
ATOM	1225	CE	LYS	156		45.685	16.692	57.005		39.79	AAAA
ATOM	1226	NZ	LYS	156		45.671	15.192	57.058		36.33	AAAA
ATOM	1227	C	LYS	156		45.233	17.260	62.299		26.40	AAAA
ATOM	1228	Ö	LYS	156		46.188	16.511	62.529		26.19	AAAA
ATOM	1229	N	LYS	157		44.960	18.337	63.032		22.50	AAAA
ATOM	1230	CA	LYS	157		45.757	18.709	64.204		21.12	AAAA
MOTA	1231	CB	LYS	157		45.535	20.181	64.591		28.95	AAAA
ATOM	1232	CG	LYS	157		46.160	21.215	63.652		25.94	AAAA
ATOM	1233	CD	LYS	157		47.669	21.067	63.575		35.16	AAAA
	1234	CE								39.24	AAAA
MOTA		NZ	LYS	157		48.281	22.099 21.869	62.627		40.01	AAAA
ATOM	1235		LYS	157		49.742		62.406			
MOTA	1236	C	LYS	157		45.421	17.825	65.411		22.98	AAAA
ATCM	1237	0	LYS	157		46.085	17.903	66.444		27.77	AAAA
ATCM	1238	N	GLY	158		44.392	16.995	65.284		26.49	AAAA
ATOM	1239	CA	GLY	158		44.023	16.106	66.376		24.82	AAAA
ATCM	1240	Ç	GLY	158		42.771	16.459	67.161		33.13	AAAA
ATOM	1241	0	GLY	158		42.421	15.775	68.128		27.21	AAAA
MOTA	1242	N	PHE	159		42.085	17.529	66.781		27.47	AAAA
ATOM	1243	CA	PHE	159		40.866	17.861	67.502	1.00	24.15	AAAA
ATOM	1244	CB	PHE	159		40.410	19.285	67.186	1.00	27.53	AAAA
MOTA	1245	CG	PHE	159		41.264	20.343	67.827	1.00	27.26	AAAA
ATOM	1246	CD1	PHE	159		42.439	20.785	67.220	1.00	28.12	AAAA
ATOM	1247		PHE	159		40.926	20.842	69.076	1.00	21.10	AAAA
ATOM	1248		PHE	159		43.264	21.714	67.866		26.24	AAAA
ATOM	1249	CE2	PHE	159		41.738	21.768	69.736		26.07	AAAA
ATOM	1250	cz	PHE	159		42.907	22.205	69.135		23.91	AAAA
ATOM	1251	C	PHE	159		39.792	16.854	67.120		28.02	AAAA
	1252	ō	PHE	159		39.639	16.533	65.947		21.14	AAAA
ATCM				160		39.056		68.110		24.79	AAAA
ATOM	1253	N	LYS				16.361			24.75	
ATOM	1254	CA	LYS	160		38.011	15.366	67.881	1.00	24.20	AAAA

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	•							
	TOM 1255	_	YS 160	38.3	360 14.0	98 68.668	1 00 22 05	
	TOM 1256		YS 160	39.6				
	TOM 1257	_	YS 160	40.2	_			
	TOM 1258		YS 160	39.2			00 33.03	
	TOM 1259	_	YS 160	38.1			1.00 62.87	
	TOM- 1260		YS 160	36.5			1.00 21.12	
	POM 1261		/S 160	35.6				AAAA
	TOM 1262			36.4	76 17.04		1.00 19.68	AAAA
A.		CA AI		35.1			1.00 20.84	AAAA
	TOM 1264	CB A		34.8			1.00 26.02	AAAA
· A?		CG AF		34.7		71.080		AAAA
	POM 1266	CD AF		34.2			1.00 30.38	AAAA
	TOM 1267	NE AF		35.0	98 16.73	4 73.445	1.00 32.99	AAAA
	OM 1268	CZ AF		36.2	72 16.27	8 73.883	1.00 40.49	AAAA
λΊ		NH1 AR	,	36.7			1.00 31.49	AAAA
	OM 1270	NH2 AR		37.00	03 17.01	4 74.712	1.00 38.54	AAAA
	OM 1271	C AR		35.13	71 19.06	0 68.680	1.00 18.98	AAAA -
	OM 1272	O AR		35.55	52 19.93		1.00 23.57	AAAA
	OM 1273	N IL		34.74		2 67.458	1.00 19.82	AAAA AAAA
	OM 1274 OM 1275	CA IL		34.74		0 66.947	1.00 17.81	AAAA
	OM 1275 OM 1276	CB IL		35.52		7 65.626	1.00 18.33	AAAA
AT		CG2 IL		35.54			1.00 13.65	AAAA
AT		CG1 ILI		36.93			1.00 18.15	AAAA
ATY		CD1 ILI	. –	37.72		2 64.670	-1.00 22.52	AAAA
ATY		C ILI		33.31			1.00 14.71	AAAA
ATY		N LEU		32.52	_		1.00 17.99	AAAA
AT		CA LET		32.99	_		1.00 16.93	AAAA
ATO		CB LEU		31.65			1.00 20.73	AAAA
ATC		CG LEU		31.11			1.00 18.45	AAAA
ATC		CD1 LEU		. 29.84			1.00 19.99	AAAA
ATC		CD2 LEU		28.65° 29.60			1.00 15.66	AAAA
ATC	M 1287	C LEU	163	31.70	5 24.751 5 24.071		1.00 18.74	AAAA
ATC	M 1288	O LEU	163	32.607			1.00 18.40	AAAA
ATC	M 1289	N TYR	164	30.752	24.128	66.188 65.186	1.00 18.65	AAAA
ATC		CA TYR	164	30.656	25.246		1.00 16.97.	AAAA
ATO		CB TYR	164	30.782	24.754		1.00 11.76 1.00 14.07	AĄĄĄ
ATO	_	CG TYR	164 .	30.593			1.00 14.07	AAAA
ATO		CD1 TYR	164	31.573		61.562	1.00 27.08	AAAA
ATO		CE1 TYR	164	31.353	27.832	60.598	1.00 26.21	AAAA
ATO		CD2 TYR	164	29.415		61.070	1.00 21.45	AAAA AAAA
ATO		CE2 TYR	164	29.193	26.891	60.137	1.00 21.89	AAAA
ATO		Z TYR	164	30.148		59.896	1.00 16.35	AAAA
ATO ATO		OH TYR	164	29.857	28.764	58.913	1.00 27.44	AAAA
ATO		TYR	164	29.279		64.463	1.00 15.67	AAAA
ATO			164	28.^60		64.455	1.00 16.07	AAAA
ATO		A ILE	165	2940	27.187	64.674	1.00 14.52	AAAA
ATO		B ILE	165	27.178	27.887	64.893	1.00 18.37	AAAA
ATOM		G2 ILE	165 165	27.959	28.596		1.00 13.31	AAAA
ATOM	. "::.:: -	G1 ILE	165	26.654 28.172	29.359	66.419	1.00 13.06	AAAA
ATOM		D1 ILE	165	28.493	27.573		1.00 17.28	AAAA
ATOM			165	27.853	28.209	68.739	1.00 15.02	AAAA
ATOM			165	28.759	28.926	63.779	1.00 20.75	AAAA
ATOM			166		29.733 28.901	63.569	1.00 16.67	AAAA
ATOM			166	26.503	29.779		1.00 15.37	AAAA
ATOM	1311 CI		166	26.276	28.885		1.00 15.63	AAAA
ATOM	1312 C		166	26.279	29.666		1.00 12.31	AAAA
ATOM	1313 OI	Ol ASP	166	25.378	30.508		1.00 16.37	AAAA
ATOM	1314 OF	2 ASP	166	27.187	29.428		1.00 13.41	AAAA
ATOM	1315 C	ASP	166	25.334	30.740		1.00 16.06	AAAA
ATOM	1316 🤉	ASP	166	24.160	30.740		1.00 15.54	AAAA
MOTA	1317 N	LEU	167	25.647	32.010		00 12.60	AAAA
ATOM	· 1318 CA		167	24.598	32.993		00 14.02	AAAA
ATOM	1319 ÇB		167	25.051	33.962		.00 12.05	AAAA
ÄTCM	1320 CG	LEU	167	25.345	33.239	CO	.00 14.61	AAAA
.•	•	•				-2031 1	.00 17.20	AAAA
								•

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ATOM	1321	CD	LEU	167	25.635	34.271	66.169	1.00 28.82	AAAA
ATOM	1322	CD2	LEU	167	24.148	32.372	65.513	1.00 18.59	AAAA
ATOM	1323	С	LEU		24.122	33.776	61.449	1.00 12.62	AAAA
	1324	ō	LEU	167	23.288	34.678	61.570	1.00 15.00	AAAA
MOTA	1325	N	ASP	168	24.667	33.431	60.288	1.00 14.35	AAAA
ATOM					24.277	34.056	59.022	1.00 19.50	AAAA
ATOM	1326	CA	ASP			33.409	57.880	1.00 25.15	
MOTA	1327	CB	ASP	168	25.060				AAAA
MOTA	1328	CG	ASP	168	24.908	34.145	56.573	1.00 48.45	AAAA
MOTA	1329		ASP	168	25.477	35.247	56.454	1.00 64.45	AAAA
MOTA	1330	OD2	2 ASP	168	24.215	33.633	55.668	1.00 44.71	AAAA
MOTA	1331	С	ASP	168	22.787	33.751	58.834	1.00 16.30	AAAA
MOTA	1332	0	ASP	168	22.327	32.696	59.252	1.00 17.72	AAAA
ATOM	1333	N	ALA	169	22.059	34.657	58.175	1.00 14.11	AAAA .
MOTA'	1334	CA	ALA	169	20.618	34.503	57.934	1.00 19.61	AAAA
ATOM	1335	CB	ALA	169	20.006	35.856	57.470	1.00 13.56	AAAA
ATOM	1336	С	ALA	169	20.277	33.400	56.926	1.00 18.23	AAAA
ATOM	1337	0	ALA	169	19.105	33.159	56.641	1.00 17.20	AAAA
ATOM	1338	N.	HIS	170	21.301	32.750	56.373	1.00 16.53	AAAA
ATOM	1339	CA	HIS	170	21.075	31.652	55.436	1.00 17.51	AAAA
	1340	CB	HIS	170	21.616	31.973	54.033	1.00 22.32	AAAA
MOTA		CG	HIS	170	20.954	33.142	53.377	1.00 25.38	AAAA
ATOM	1341				19.934	33.196	52.487	1.00 19.33	AAAA
ATOM	1342		HIS	170				1.00 13.33	AAAA
ATOM	1343		HIS	170	21.308	34.448	53.638		
MOTA	1344		HIS	170	20.535	35.257	52.935	1.00 30.34	AAAA
ATOM	1345		HIS	170	19.692	34.523	52.229	1.00 17.51	AAAA
ATOM	1346	С	HIS	170	21.781	30.413	55.967	1.00 16.72	AAAA .
MOTA	1347	0	HIS	170	22.827	30.511	56.610	1.00 15.92	AAAA
ATOM	1348	N	HIS	171	21.209	29.245	55.682	1.00 15.28	AAAA
MOTA	1349	CA.	HIS	171	21.751	27.961	56.123	1.00 12.53	AAAA
MOTA	1350	CB	HIS	171	20.702	26.878	55.814	1.00 14.09	AAAA
MOTA	1351	CG	HIS	171	21.180	25.468	55.980	1.00 17.27	AAAA
MOTA	1352	CD2	HIS	171	21.249	24.447	55.090	1.00 12.48	AAAA
ATOM	1353	ND1	HIS	171	21.622	24.956	57.181	1.00 26.73	AAAA
MOTA	1354	CE1	HIS	171 ·	21.948	23.685	57.021	1.00 15.98	AAAA
ATOM	1355	NE2	HIS	171	21.729	23.352	55.761	1.00 20.03	AAAA
ATOM	1356	С	HIS	171	23.107	27.602	55.498	1.00 15.55	AAAA
ATOM	1357	0	HIS	171	23.318	27.784	54.298	1.00 17.03	AAAA
ATOM	1358	N	CYS	172	24.026	27.105	56.323	1.00 14.33	AAAA
ATOM	1359	CA	CYS	172	25.350	26.675	55.866	1.00 13.65	AAAA
ATOM	1360	CB	CYS	172	26.330	26.631	57.054	1.00 12.99	AAAA
ATOM	1361	SG	CYS	172	25.680	25.826	58.551	1.00 17.17	AAAA
ATOM	1362	c	CYS	172	25.212	25.274	55.257	1.00 16.52	AAAA
ATOM	1363	ŏ	CYS	172	25.750	24.297	55.783	1.00 14.95	AAAA
	1364	N	ASP	173	24.516	25.173	54.130	1.00 15.42	AAAA
ATOM	1365	CA	ASP	173	24.302	23.865	53.531	1.00 14.75	AAAA
ATOM	1366	CB	ASP	173	23.339	23.956	52.332	1.00 17.73	AAAA
ATOM		CG	ASP	173	23.765	24.966	51.283	1.00 22.84	AAAA
ATOM	1367				23.106	24.998	50.216	1.00 18.68	AAAA
MOTA	1368		ASP	173	24.730	25.728	51.504	1.00 15.34	AAAA
ATOM	1369		ASP	173		23.726	53.149	1.00 16.39	AAAA
MOTA	1370	C	ASP	173	25.590		53.279	1.00 16.48	AAAA
ATOM	1371	0	ASP	173	25.684	21.922		1.00 15.58	AAAA
MOTA	1372	N	GLY	174	26.583	23.912	52.705		AAAA
MOTA	1373	CA	GLY	174	27.869	23.346	52.360	1.00 13.97	AAAA
ATOM	1374	C	GLY	174	28.508	22.723	53.595	1.00 18.44	
ATOM	1375	0	GLY	174	28.970	21.586	53.540	1.00 15.48	AAAA
ATOM	1376	N	VAL	175	28.554	23.456	54.706	1.00 16.84	AAAA
ATOM	1377	CA	AYT	175	29.136	22.923	55.946	1.00 16.54	AAAA
ATOM	1378	CB	VAL	175	29.201	24.031	57.037	1.00 15.88	AAAA
ATOM	1379		VAL	175	29.927	23.507	58.307	1.00 15.35	AAAA
ATOM	1380		VAL	175	29.923	25.258	56.476	1.00 15.62	AAAA
ATOM	1381	c	VAL	175	28.318	21.720	56.467	1.00 19.21	AAAA
ATOM	1382	ō	VAL	175	28.876	20.735	56.961	1.00 17.75	AAAA
ATOM	1383	N	GLN	176	26.996	21.798	56.367	1.00 17.74	AAAA
	1384	CA	GLN	176	26.164	20.685	56.832	1.00 15.66	AAAA
ATOM	1385	CB	GLN	176	24.678	20.973	56.595	1.00 16.64	AAAA
MOTA		CG	GLN	176	23.789	19.788	56.952	1.00 17.00	AAAA
MOTA	1386	٠.	7سوب		23.,05	,00	-		•

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MOTA	1387	CD GLN	176	22.325	20.106	56.884	1.00 21.52	AAAA
MOTA	1388		176	21.850	21.016	57.567	1.00 21.72	AAAA
MOTA	1389	NE2 GLN	176	21.581	19.348	56.064	1.00 20.30	AAAA
MOTA	1390	C GLN	176	26.527	19.387	56.121	1.00 16.33	AAAA
MOTA	1391	O GLN	176	26.751	18.354	56.748	1.00 17.46	AAAA
MOTA	1392	N GLU	177	26.581	19.443	54.799	1.00 22.24	AAAA
ATOM	1393	CA GLU	177	26.909	18.251	54.021	1.00 19.67	AAAA
MOTA	1394	CB GLU	177	26.857	18.587	52.533	1.00 15.55	AAAA
MOTA	1395	CG GLU	177	27.131	17.388	51.623	1.00 20.24	AAAA
•	_							
MOTA	1396	CD GLU	177	26.960	17.740	50.159	1.00 27.00	AAAA
MOTA	1397	OE1 GLU	177	27.974	17.935	49.450	1.00 30.23	AAAA
ATOM	1398	OE2 GLU	177	25.796	17.853	49.725	1.00 26.89	AAAA
ATOM	1399	C GLU	177	28.284	17.713	54.376	1.00 20.42	AAAA
MOTA	1400	O GLU	177	28.486	16.503	54.527	1.00 17.05	AAAA
ATOM	1401	N ALA	178	29.233	18.626	54.527	1.00 19.67	AAAA
ATOM	1402	CA ALA	178	30.611	18.259	54.839	1.00 18.18	AAAA
ATOM	1403	CB ALA	178	31.464	19.519	54.918	1.00 12.76	AAAA
MOTA	1404	C ALA	178	30.806	17.418	56.106	1.00 17.56	AAAA
ATOM	1405	O ALA	178	31.690	16.555	56.167	1.00 17.72	AAAA
ATOM	14.06	N . PHE.	179	29.981	17.656	57.116	1.00 18.82	AAAA
ATOM	1407	CA PHE	179	30.124	16.945	58.379	1.00 20.26	AAAA
			179	30.554	17.948			
MOTA	1408	CB PHE				59.439	1.00 13.17	AAAA
ATOM	1409	CG PHE	179	31.779	18.693	59.048	1.00 16.28	AAAA
		CD1 PHE	179	31.705	20.017		1.00 13.77	
ATOM	1410					58.610		AAAA
ATOM	1411	CD2 PHE	179	33.002	18.031	58.995	1.00 15.57	AAAA
	1412	CE1 PHE	179	32.845	20.673	58.114	1.00 20.03	AAAA
ATOM	-							
ATOM	1413	CE2 PHE	179	34.145	18.677	58.500	1.00 20.30	AAAA
ATOM	1414	CZ PHE	179	34.060	20.002	58.058	1.00 19.51	AAAA
MOTA.	1415	C PHE	179	28.882	16.219	58.833	1.00 18.52	AAAA
MOTA	1416	O PHE	179	28.773	15.828	60.000	1.00 20.21	AAAA
MOTA	1417	n tyr	180	27.969	16.016	57.895	1.00 18.33	AAAA
MOTA	1418	CA TYR	180	26.698	15.379	58.176	1.00 19.93	AAAA
	1419		180	25.874	15.310		1.00 20.97	AAAA
ATOM								
MOTA	1420	CG TYR	180	24.402	15.341	57.159	1.00 19.80	AAAA
ATOM	1421	CD1 TYR	180 -	23.565	14.337	56.686	1.00 23.87	AAAA
ATOM	1422	CE1 TYR	180	22.203	14.391	56.898	1.00 21.32	AAAA
ATOM	1423	CD2 TYR	180	23.831	16.416	57.865	1.00 19.02	AAAA
ATOM	1424	CE2 TYR	180	22.470	16.482	58.084	1.00 26.84	AAAA
ATOM	1425	CZ TYR	180	21.659	15.462	57.594	1.00 30.54	AAAA
				20.310			1.00 22.81	AAAA
MOTA	1426		180		15.514	57.794		
ATOM	1427	C TYR	180	26.855	13.970	58.737	1.00 22.61	AAAA
	1428	O TYR	180	26.064	13.526	59.579	1.00 23.44	AAAA
· ATOM								
ATOM	1429	N ASP	181	27.893	13.298	58.253	1.00 22.27	AAAA
ATOM	1430	CA ASP	181	28.245	11.920	58.590	1.00 33.84	AAAA
MOTA	1431	CB ASP	181	28.916	11.318	57.339	1.00 41.74	AAAA
MOTA	1432	CG ASP	181	30.035	10.363	57.662	1.00 57.71	AAAA
				30.999		58.340	1.00 61.40	AAAA
MOTA	1433	OD1 ASP	181					
ATOM	1434	OD2 ASP	181	29.965	9.197	57.221	1.00 65.77	AAAA
	1435	C ASP	181	29.107	11.654	59.838	1.00 30.21	AAAA
MOTA								
MOTA	1436	O ASP	181	29.307	10.497	60.227	1.00 27.84	AAAA
	1437	N THR	182	29.615	12.696	60.480	1.00 27.53	AAAA
ATOM								
ATOM	1438	CA THR	182	30.472	12.466	61.634	1.00 21.19	AAAA
ATOM	1439	CB THR	182	31.918	12.977	61.358	1.00 26.55	AAAA
MOTA	1440	OG1 THR	182	32.729	12.763	62.513	1.00 25.62	AAAA
ATOM	1441	CG2 THR	182	31.922	14.471	61.037	1.00 21.67	AAAA
ATOM	1442	C THR	182	30.010	13.050	62.954	1.00 25.02	AAAA
ATOM	1443	O THR	182	29.306	14.049	62.992	1.00 23.56	AAAA
							1.00 19.66	AAAA
MOTA	1444	N ASP	183	30.434	12.424	64.042		
ATOM	1445	CA ASP	183	30.086	12.894	65.371	1.00 21.52	AAAA
				29.735			1.00 28.52	AAAA
ATOM	1446	CB ASP	183		11.700	66.275		
ATOM	1447	CG ASP	183	30.920	10.783	66.523	100 32.30	AAAA
	1448		183	31.667	10.502	65.565	1.00 30.99	AAAA
MOTA		OD1 ASP						
MOTA	1449	OD2 ASP	183	31.095	10.326	67.675	1.00 48.65	AAAA
	1450	C ASP	183	31.257	13.685	65.947	1.00 16.66	AAAA
ATOM								
ATOM	1451	O ASP	183	31.236	14.092	67.104.	1.00 23.37	AAAA
MOTA	1452	N GLN	184	32.286	13.909	65.131	1.00 21.95	AAAA
PION.	¥ 3 3 6	., GLIN	-04	22.000				
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Figure 16-23

		2.452			204		4			
	ATOM	1453	CA	GLN	184	33.437	14.672	65.590	1.00 17.65	AAAA
	ATOM	1454	CB	GLN	184	34.701	14.243	64.866	1.00 21.36	AAAA
		1455								
	MOTA			GLN	184	35.068		65.102	1.00 27.38	AAAA
	ATOM	1456	CD	GLN	184	36.485	12.476	64.691	1.00 31.96	AAAA
	MOTA	1457	OE:	1 GLN	184	36.899		63.573	1.00 29.90	
										AAAA
	ATOM	1458	NE.	2 GLN	184	37.239	11878	65.599	1.00 31.84	AAAA
•	ATOM	1459	С	GLN	184	33.207	16.165	65.382	1.00 18.54	AAAA
	ATOM	1460	ō	GLN	184					
						33.881	17.009	65.972	1.00 18.11	AAAA
	MOTA	1461	N	VAL	185	32.258	16.481	64.519	1.00 19.18	AAAA
	MOTA	1462	CA	VAL	185	31.934	17.872	64.267	1.00 21.57	AAAA
	ATOM	1463	CB	VAL	185	. 32.261	18.264	62.807	1.00 22.64	AAAA
	ATOM	1464	CG:	LVAL	185	31.994	19.768	62.591	1.00 16.26	AAAA
	MOTA	1465	CG	VAL	185	33,722	17.924	62.500	1.00 16.77	ÀAAA
	ATOM	1466	С	VAL	185	30.449	18.035	64.523	1.00 15.91	AAAA
	ATOM	1467	0	VAL	185	29.658	17.156	64.179	1.00 20.79	AAAA
		1468	N	PHE			19.146			
	ATOM					30.081		65.153	1.00 18.73	AAAA
	ATOM	1469	CA	PHE	186	28.687	19.446	65.435	1.00 16.22	AAAA
	MOTA	1470	CB	PHE	186	28.432	19.559	66.952	1.00 16.83	AAAA
	. ATOM	1471	CG	PHE	186	26.976	19.682	67.299	1.00 17.96	. AAAA
	MOTA	1472	CDI	PHE	· 186	26.319	18.656	67.968	1.00 23.24	AAAA
	ATOM	1473	CD2	PHE	186	26.240	20.797	66.904	1.00 15.41	AAAA
		1474		PHE	186	24.953	18.738			
	ATOM							68.235	1.00 18.99	AAAA
	MOTA	1475	CE2	PHE	186	24.879	20.887	67.168	1.00 24.05	AAAA
	MOTA	1476	CZ	PHE	186	24.234	19.846	67.838	1.00 22.93	AAAA
	ATOM	1477	C	PHE	186	28.437	20.789	64.778	1.00 17.16	AAAA
	ATOM	1478	0	PHE	186	29.192	21.725	64.993	1.00 19.37	AAAA
	ATOM	1479	N	VAL	187	27.391	20.874	63.961	1.00 19.67	AAAA
	MOTA	1480	CA	VAL	187	27.075	22.116	63.277	1.00 17.74	AAAA
		1481								
	ATOM		CB	VAL	187	27.010	21.914	61.720	1.00 18.65	AAAA
	MOTA	1482	CG1	VAL	` 187	26.578	23.211	61.024	1.00 17.31	AAAA
	ATOM	1483	CG2	VAL	187	28.359	21.453	61.194	1.00 16.65	AAAA
	ATOM	1484	С	VAL	187	25.732	22.637	63.746	1.00 18.46	AAAA
	MOTA	1485	0	VAL	187	24.752	21.903	63.764	1.00 20.64	AAAA
	ATOM	1486	Ņ	LEU	188	25.708	23.899	64.150	1.00 14.42	AAAA
	ATOM	1487	CA	LEU	188	24.482	24.563	64.567	1.00 16.68	AAAA
	ATOM	1488	CB	LEU	188	24.568	25.070	66.009	1.00 13.98	AAAA
	MOTA	1489	CG	LEU	188	23.522	26.119	66.450	1.00 13.66	AAAA .
	ATOM	1490	CD1	LEU	188	22.103	25.556	66.401	1.00 15.55	AAAA
	ATOM	1491	CD2	LEU	188	23.844	26.585	67.861	1.00 16.40	AAAA
		1492			188					
	MOTA		С	LEU		24.272	25.756	63.667	1.00 20.01	AAAA
	ATOM	1493	0	LEU	188	25.164	26.595	63.506	1.00 18.86	AAAA
	ATOM	1494	N	SER	189	23.106	25.845	63.057	1.00 14.46	AAAA
	ATOM	1495	CA	SER	189	22.841	27.011	62.230	1.00 14.56	AAAA
	ATOM	1496	CB	SER	189	22.896	26.668	60.737	1.00 15.55	AAAA
	ATOM	1497	OG	SER	189	22.619	27.851	60.008	1.00 14.09	AAAA
	MOTA	1498	Ċ	SER	189	21.487	27.606	62.508	1.00 15.24	AAAA
_	MOTA	1499	0	SER	189	20.509	26.885	62.578	1.00 21.46	AAAA
	ATOM	1500	N	LEU	190	21.423	28.921	62.690	1.00 14.92	AAAA
	MOTA	1501	CA	LEU	190	20.128	29.572	62.826	1.00 15.54	AAAA
		1502		LEU		20.084				
	MOTA		CB		190		30.663	63.906	1.00 21.02	AAAA
	ATOM	1503	CG	LEU	190	20.594	30.532	65.339	1.00 30.17	AAAA
	ATOM	1504	CD1	LEU	190	19.736	31.437	66.210	1.00 19.75	AAAA
		1505	CD2		190	20.547	29.130	65.831	1.00 19.08	AAAA
	ATOM									
	ATOM	1506	С	LEU	190 ·	20.035	30.250	61.456	1.00 14.31	AAAA
	MOTA	1507	0	LEU	190	21.031	30.752	60.951	1.00 15.43	AAAA
	ATOM	1508	Ν.	HIS	191	18.855	30.285	60.856	1.00 16.88	AAAA
	MOTA	1509	CA	HIS	191	18.732	30.884	59.535	1.00 14.34	AAAA
	ATOM	1510	CB	HIS	191	19.506	30.015	58.539	1.00 17.34	AAAA
	ATOM	1511	CG	HIS	191	19.229	28.546	58.697	1.00 14.27	AAAA
					191	19.941				
	ATOM	1512	CD2				27.578	59.319	1.00 9.60	AAAA
	ATOM	1513	ND1		191	18.073	27.940	58.247	1.00 21.22	AAAA
	ATOM	1514	CE1	HIS	191	18.088	26.660	58.582	1.00 17.22	AAAA
		1515	NE2		191	19.212	26.415	59.232	1.00 20.70	AAAA
	ATOM									
	ATOM	1516	С	HIS	191	17.277	31.026	59.110	1.00 16.19	AAAA
	ATOM	1517	0	HIS	191	16.381	30.489	59.766	1.00 16.73	AAAA
		- 1518	N	GLN	192	17.044	31.796	58.045	1.00 14.78	AAAA
	A1011.						54.,50	-		
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Figure 16-24

				-					
MOTA	151	9 C	A GLN	192	15.683	31.968	57.516	1.00 16.33	KAAA
ATOM	152	0 C	B GLN		15.669				
ATOM	152								AAAA
					16.174				AAAA
MOTA	152	_			16.408				AAAA
MOTA	152		El GLN	192	15.490	35.566	54.587	1.00 20.46	AAAA
ATOM	. 152	4 NI	E2 GLN	192	17.630	34.839	54.665		AAAA
ATOM	152	5 C	GLN	192	15.262				
ATOM	152		GLN		16.071				AAAA
	152								AAAA
AŢOM			SER		14.007			1.00 15.63	AAAA
ATOM	152		-		13.561	28.907	56.877	1.00 13.84	AAAA
ATOM	1529	9 CE	SER	193	12.097	28.677	57.284	1.00 17.28	AAAA
ATOM	1530	000	SER	193	11.639	27.439		1.00 17.58	
ATOM	1531	L C	SER		- 13.687			-1.00 11.80	AAAA
ATOM	1532		SER						AAAA
								1.00 18.44	AAAA
MOTA	1533		PRO		. 14.103			1.00 14.59	AAAA
ATOM	1534			194	14.335	26.325	55.782	1.00 19.22	AAAA
MOTA	1535	5 CA	PRO	194	14.268	27.143	53.513	1.00 15.30	AAAA
ATOM	153€	CB	PRO	194	14.892			1.00 18.33	
ATOM	1537			194	15.359				AAAA
ATOM	1538							1.00 22.34	AAAA
			PRO	194	12.880			1.00 16.40	AAAA
MOTA	1539		PRO	194	12.757	27.003		1.00 19.43	AAAA
ATOM	1540	N	GLU	195	11.828	27.151	53.681	1.00 20.57	AAAA
ATOM	1541	. CA	GLU	195	10.483	27.161	53.099	1.00 30.15	AAAA
ATOM	1542	CB	GLU	195	9.386	27.037		1.00 31.91	
ATOM	1543		GLU	195	8.987	28.325			
ATOM	1544			195				1.00 45.60	AAAA
	1545				7.880	29.119		1.00 34.45	AAAA
ATOM			1 GLU	195	7.635	30.259		1.00 43.98	Aaaa
ATOM	1546		2 GLU	195	. 7.241	28.627		1.00 38.39	AAAA
ATOM	1547	С	GLU	195	10.333	28.474	52.318	1.00 26.92	AAAA
ATOM	1548	0	GLU	195	9.522	28.557		1.00 24.59	AAAA
MOTA	1549	N	TYR	196	11.116	29.501	52.669	1.00 18.16	AAAA
MOTA	1550	CA	TYR	196	11.024	30.753	51.922		
ATOM ·	1551	CB	TYR	196				1.00 15.81	AAAA
					10.208	31.801	52.690	1.00 20.01	AAAA
ATOM	1552	CG	TYR	196	10.868	32.353	53.932	1.00 19.77	AAAA
ATOM	1553		LTYR	196	11.779	33.408	53.853	1.00 18.24	AAAA
ATOM	1554	CEI	L TYR	196	12.407	33.898	54.988	1.00 18.50	AAAA
ATOM	1555	CD2	TYR	196	10.598	31.801	55.185	1.00 18.12	AAAA
ATOM	1556	CE2	TYR	196	11.223	32.283	56.339	1.00 21.09	AAAA
MOTA	1557	CZ	TYR	196	12.125	33.326	56.235	1.00 20.39	
ATOM	1558	ОН	TYR	196	12.759				AAAA
	1559					33.784	57.367	1.00 16.20	AAAA
ATOM		Ç	TYR	196	12.342	31.372	51.475	1.00 16.89	AAAA
ATOM	1560	0	TYR	196	12.336	32.347	50.718	1.00 23.08	AAAA
ATOM	1561	N	ALA	197	13.466	30.817	51.911	1.00 17.52	AAAA
ATOM	1562	CA	ALA	197	14.754	31.400	51.512	1.00 20.26	AAAA
ATOM	1563	CB	ALA	197	15.315	32.261	52.659	1.00 20.74	AAAA
ATOM	1564	С	ALA	197	15.814	30.392	51.074	1.70 13.51	
ATOM	1565	ō	ALA	197	15.787				AAAA
ATOM	1566	N				29.229	51.457	1 30 19.35	AAAA
			PHE	198	16.757	30.869	50.257	1. 70 18.01	AAAA
ATOM	1567	CA	PHE	198	17.861	30.049	49.782	1.00 17.97	AAAA
ATOM	1568	CB	PHE	198	18.929	30.933	49.119	1.00 20.38	AAAA
ATOM	1569	CG	PHE	198	20.094	30.162	48.545	1.00 23.61	AAAA
ATOM	1570	CD1	PHE	198	20.039	29.660	47.245	1.00 29.71	AAAA
MOTA	1571		PHE	198	21.229	29.899			
ATOM	1572						49.321	1.00 19.06	AAAA
			PHE	198	21.091	28.908	46.719	1.00 30.39	AAAA
ATOM	1573	CE2		198	22.290	29.145	48.807	1.00 23.17	AAAA
MOTA	1574	CZ	PHE	198	22.218	28.646	47.493	1.00 22.74	AAAA
MOTA	1575	2	PHE	198	18.453	29.419	51.032	1.00 16.02	AAAA
ATOM	1576	0	PHE	198	18.552	30.073	52.061	1.00 20.95	AAAA
ATOM	1577	N	PRO	199	18.941				
						28.176	50.937	1.00 19.92	AAAA
ATOM	1578	CD	PRO	199	19.600	27.508	52.074	1.00 17.86	AAAA
ATOM	1579	CA	PRO	199	18.990	27.318	49.744	1.00 23.54	AAAA
MOTA	1580	CB	PRO	199	20.108	26.344	50.095	1.00 22.70	AAAA
MOTA	1581	CG	PRO	199	19.813	26.087	51.534	1.00 23.16	AAAA
	1582	c	PRO	199	17.710	26.595	49.312	1.00 30.97	AAAA
ATOM	1583	0	PRO	199	17.733				
						25.855	48.322	1.00 23.25	AAAA
ATOM	1584	N	PHE	200	16.621	26.795	50.054	1.00 20.32	AAAA

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					·				
ATOM	1585	CA	PHE	200	15.319	26.166	49.752	1.00 20.27	AAAA
	-		PHE	200	14.840	26.533	48.346	1.00 19.77	AAAA
ATOM	1586	CB	-						
ATOM	1587	CG	PHE	200	14.752	27.999	48.082	1.00 18.06	AAAA
MOTA	1588	CD1	PHE	200	15.742	28.644	47.346	1.00 18.97	AAAA
			PHE	200	13.654	28.736	48.519	1.00 19.06	AAAA
MOTA	1589								
ATOM	1590	CEl	PHE	200	15.635	30.003	47.042	1.00 21.67	AAAA
MOTA	1591	CE2	PHE	200	13.539	30.101	48.221	1.00 22.60	AAAA
	1592	cz	PHE	200	14.527	30.736	47.482	1.00 18.93	AAAA
ATOM									
ATOM	1593	С	PHE	200	15.294	24.637	49.845	1.00 18.44	AAAA
ATOM	1594	0	PHE	200	14.302	24.049	50.272	1.00 20.74	AAAA
	1595	N	GLU	201	16.384	24.004	49.418	1.00 20.77	AAAA
MOTA									
MOTA	1596	CA	GLU	201	16.522	22.542	49.399	1.00 27.34	AAAA
MOTA	1597	CB	GLU	201	17.498	22.146	48.284	1.00 28.99	AAAA ·
ATOM	1598	CG	GLU	201	17.024	22.458	46.881	1.00 34.82	AAAA
							45.848	1.00 32.40	
ATOM	1599	CD	GLU	201	18.123	22.265			AAAA
MOTA	1600	OE1	GLU	201	18.701	21.155	45.769	1.00 38.28	AAAA
ATOM	1601	OF2	GLU	201	18.405	23.230	45.111	1.00 40.08	AAAA
				201	17.007	21.891	50.695	1.00 23.51	AAAA
ATOM	1602	C	GLU						
ATOM	1603	0	GLU	201	16.845	20.689	50.886	1.00 23.17	AAAA
MOTA	1604	N	LYS	202	17.619	22.681	51.571	1.00 20.03	AAAA
	1605	CA	LYS	202	18.178	22.177	52.829	1.00 17.01	AAAA
MOTA									
MOTA	1606	CB	LYS	202	19.666	21.862	52.634	1.00 19.24	AAAA
MOTA	1607	CG	LYS	202	19.903	20.769	51.611	1.00 36.04	AAAA
ATOM	1608	CD	LYS	202	20.997	21.162	50.648	1.00 45.11	AAAA
		CE	LYS	202	21.060	20.209	49.463	1.00 55.83	AAAA
MOTA	1609							1.00 28.09	
ATOM	1610	NZ	LYS	202	22.024	20.662	48.422		AAAA
ATOM	1611	С	LYS	202	18.016	23.240	53.899	1.00 17.02	AAAA
ATOM	1612	0	LYS	202	. 17.705	24.381	53.585	1.00 20.20	AAAA
				203	18.232	22.875	55.160	1.00 22.94	AAAA
MOTA	1613	N	GLY						
MOTA	1614	CA	GLY	203	18.064	23.850	56.223	1.00 19.38	AAAA
MOTA	1615	С	GLY	203	16.874	23.564	57.128	1.00 20.48	AAAA
MOTA	1616	0	GLY	203	16.607	24.312	58.070	1.00 18.55	AAAA
				204	16.150	22.484	56.852	1.00 15.42	AAAA
ATOM	1617	N	PHE						
MOTA	1618	CA	PHE	204	14.983	22.143	57.670	1.00 20.73	AAAA
MOTA	1619	CB	PHE	204	14.018	21.212	56.903	1.00 19.97	AAAA
ATOM	1620	CG	PHE	204	13.441	21.838	55.667	1.00 19.63	AAAA
		CD1		204	14.137	21.801	54.459	1.00 24.96	AAAA
MOTA	1621						55.725	1.00 18.92	AAAA
ATOM	1622	CD2	PHE	204	12.230	22.523			
MOTA	1623	CE1	PHE	204	13.636	22.438	53.327	1.00 20.66	AAAA
ATOM	1624	CE2	PHE	204	11.720	23.169	54.597	1.00 24.86	AAAA
	1625	cz	PHE	204	12.422	23.127	53.400	1.00 23.66	AAAA
ATOM						21.513	59.006	1.00 18.73	AAAA
ATOM	1626	С	PHE	204	15.376				
ATOM	1627	0	PHE	204	16.415	20.851	59.131	1.00 20.18	AAAA
ATOM	1628	N	LEU	205	14.518	21.726	59.994	1.00 19.46	AAAA
			LEU	205	14.727	21.244	61.356	1.00 21.09	AAAA
MOTA	1629					21.674	62.233	1.00 23.44	· AAAA
MOTA	1630		LEU	205	13.547				
ATOM	1631	CG	LEU	205	13.506	21.222	63.693	1.00 23.23	AAAA
ATOM	1632	CD1	LEU	205	14.717	21.736	64.445	1.00 24.06	AAAA
	1633	CD2		205	12.224	21.743	64.312	1.00 30.63	AAAA
MOTA	_				14.943	19.748	61.489	1.00 23.53	AAAA
ATOM	1634	С	LEU	205					
ATOM	1635	0	LEU	205	15.659	19.315	62.381	1.00 21.28	AAAA
ATOM	1636	N	GLU	206	14.356	18.959	60.591	1.00 21.59	AAAA
				206	14.487	17.502	60.686	1.00 27.89	AAAA
ATOM	1637		GLU						
ATOM ·	1638	CB	GLU	206	13.345	16.816	59.928	1.00 28.90	AAAA
MOTA	1639	CG	GLU	206	12.060	17.615	59.942	1.00 48.55	AAAA
			GLU	206	12.169	18.832	59.042	1.00 46.86	AAAA
ATOM	1640				11.360		59.178	1.00 21.58	AAAA
MOTA	1641	OE1		206		19.769		1.00 61.30	
ATOM	1642	OE2	GLU	206	13.076	18.833	58.181	1.00 63.58	AAAA
	1643		GLU	206	15.819	16.955	60.188	1.00 22.86	AAAA
MOTA				206	16 071	15.753	60.286	1.00 21.21	AAAA
MOTA	1644		GLU					1.00 25.04	AAAA
MOTA	1645		GLU	207	16.666	17.816	59.631		
ATOM	1646	CA	GLU	207	17.976	17.373	59.152	1.00 19.61	AAAA
	1647		GLU	207	18.483	18.322	58.055	1.00 20.75	AAAA
ATOM				207	17.682	18.222	56.753	1.00 18.44	AAAA
ATOM	1648		GLU						AAAA
MOTA	1649		GLU	207	17.687	19.514	55.983	1.00 24.75	
MOTA	1650	OE1	GLU	207	18.738	20.182	55.948	1.00 22.17	AAAA
				-			_		

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ATO		551	0E2 G	ւԾ 207	16.6	46 19.85	4 55.39	c :	
ATO:	M 16	552	C G	LU 207	18.9				AAAA
ATO:	M 16	553	0 G	LU 207	19.50		9 60.35		AAAA
ATO:	M 16	554		E 208	19.08				AAAA
ATO			CA II						AAAA
ATO			CB II		19.93				AAAA
ATO			CG2 II		19.11			3 1.00 28.84	AAAA
ATO					19.96			3 1.00 43.26	AAAA
			CG1 II		17.90		1 63.625	1.00 21.61	
ATO			CD1 II		17.02	9 16.16	0 64.786		AAAA
ATO			C IL		21.15				AAAA
ATO			O IL	E 208	21.78				AAAA
ATOM		62	N GL	Y 209	21.51				AAAA
ATON	1 16	63 (CA GL	Y 209	22.69				AAAA
ATOM	1 16	64 (GL GL		22.34	_			AAAA
ATOM	1 16	65 () GL		21.16				AAAA
ATOM			V GL		23.37				AAAA
ATOM			A GL						AAAA
ATOM			B GL		23.23				AAAA
ATOM					23.40				AAAA
ATOM					23.04		56.965	1.00 56.41	AAAA
		_	D GL		23.17		55.495	1.00 65 37	AAAA
ATOM			El .GLU		24.29	0 10.327	55.075	1.00 62.00	
ATOM			E2 GLU		22.15	9.894	54.768	1.00 73.24	AAAA
ATOM					24.329			1.00 31.85	AAAA
ATOM				J 210	25.447			1.00 28.85	AAAA
MOTA				211	24.012			1.00 27.84	AAAA
ATOM			A GLY	211	24.991	7.502	60.657	1.00 27.84	AAAA
MOTA	167	7 C	GLY	211	25.545		61.995	1.00 26.25	AAAA
ATOM	167	6 8	GLY	211	24.788	8.324		1.00 27.79	AAAA
MOTA	167	9 N			26.865			1.00 28.66	AAAA
ATOM	168	0 C	A LYS		27.512			1.00 34.62	AAAA .
MOTA	168	1 CI			29.029		63.393	1.00 34.39	AAAA
ATOM	168				29.505		63.273	1.00 40.40	AAAA
ATOM	168			212	29.139	6.712	62.996	1.00 53.97	AAAA
ATOM	168			212			64.131	1.00 61.93	AAAA
ATOM	168			212	29.612		63.863	1.00 62.74	AAAA
ATOM	168				31.091		63.711	1.00 70.11	AAAA
ATOM	1687		LYS	212	27.181	9.741	63.725	1.00 36.04	AAAA
ATOM	1688		LYS	212	27.109	10.126	64.897	1.00 28.34	AAAA
			GLY	213	26.959	10.543	62.688	1.00 31.47	AAAA
ATOM	1689			213	26.648	11.948	62.898	1.00 31.68	AAAA
ATOM	1690		GLY	213	25.189	12.291	63.142	1.00 28.78	AAAA
ATOM	1691		GLY	213	24.840	13.460	63.259	1.00 22.56	
MOTA	1692		LYS	214	24.317	11.292	63.222	1.00 28.54	AAAA
MOTA	1693		LYS	214	22.905	11.585	63.463	1.00 28.54	AAAA
ATOM	1694		LYS	214	22.080	10.295	63.325		AAAA
ATOM	1695	CG	LYS	214	20.583	10.461	63.224	1.00 31.03	AAAA
ATOM	1696	CD	LYS	214	19.968	9.115		1.00 38.15	AAAA
ATOM	1697	CE	LYS	214	18.490	9.220	62.844	1.00 40.49	AAAA
MOTA	1698	NZ	LYS	214	17.927		62.537	1.00 48.02	AAAA
ATOM	1699	С	LYS	214	22.834	7.924 12.160	62.064	1.00 44.99	AAAA
ATOM	1700	ō	LYS	214			64.875	1.00 26.90	AAAA
ATOM	1701	N	GLY	215	23.260	11.524	65.831	1.00 33.33	AAAA
ATOM	1702	CA	GLY		22.310	13.376	64.997	1.00 24.38	AAAA
ATOM	1703	C		215	22.230	14.034	66.290	1.00 26.03	AAAA
	1704		GLY	215	23.298	15.115	66.447	1.00 27.03	AAAA
MOTA		0	GLY	215	23.352	15.820	67.458	1.00 23.34	AAAA
MOTA	1705	N	TYR	216	24.152	15.260	65.439	1.00 22.79	AAAA
MOTA	1.706	CA	TYR	216	25.217	16.257	65.512	1.00 25.51	
MOTA	1707	CB	TYR	216	26.592	15.576		1.00 20.54	AAAA
MOTA	1708	CG	TYR	216	26.900	14.671		1.00 20.54	AAAA
MOTA	1709	CD1	TYR	216	26.221	13.464	66.757	1.00 26.47	AAAA
MOTA	1710	CE1	TYR	216	26.455	12.660		1.00 28.96	AAAA
ATOM	1711		TYR	216	27.832	15.052		1.00 33.08	AAAA
MOTA	1712	CE2	TYR	216	28.074		67.552	1.00 24.21	AAAA
ATOM	1713	CZ	TYR	216		14.254		1.00 34.45	AAAA
ATOM	1714	ОН	TYR	216	27.378	13.063		1.00 40.53	AAAA
ATOM	1715	C			27.580	12.291		1.00 45.67	AAAA
ATOM	1716		TYR	216	25.104	17.391		1.00 22.57	AAAA
AIOM	T 1 T D	0	TYR	216	26.097	18.014		1.00 19.70	AAAA
-		•		• •		-		•	

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ATOM	1717	N	ASN	217	23.889	17.635	64.027	1.00 22.88	AAAA
ATOM	1718	CA	ASN	217	23.621	18.729	63.109	1.00 22.60	AAAA
ATOM	1719	CB	ASN	217	23.453	18.240	61.671	1.00 16.61	AAAA
ATOM	1720	CG	ASN	217	23.233	19.387	60.695	1.00 17.16	AAAA
ATOM	1721	OD1		217	22.098	19.704	60.307	1.00 20.23	AAAA
MOTA	1722	ND2		217	24.320	20:032	60.309	1.00 12.18	AAAA
	1723.	C	ASN	217	22.311	19.296	63.630	1.00 17.65	AAAA
MOTA	1724	ō	ASN	217	21.381	18.550	63.894	1.00 17.63	AAAA
MOTA	1725	N	LEU	218	22.236	20.610	63.793	1.00 21.68	AAAA
ATOM	1726	CA	LEU	218	21.014	21.197	64.320	1.00 21.20	AAAA
MOTA	1727	CB	LEU	218	- 21.186	21.547	65.808	1.00 17.73	AAAA
MOTA	1728	CG	LEU	218	19.906	21.702	66.647	1.00 32.30	AAAA
MOTA		CD1		218	20.228	22.427	67.944	1.00 24.51	AAAA
MOTA	1729	CD2		218	18.862	22.464	65.903	1.00 40.08	AAAA
MOTA	1730		LEU	218	20.700	22.459	63.554	1.00 19.46	AAAA
MOTA	1731	C	LEU	218	21.467	23.425	63.615-	1.00 16.70	AAAA
MOTA	1732	0	ASN	219	19.590	22.441	62.824	1.00 15.43	AAAA
MOTA	1733	N	ASN	219	19.143	23.609	62.072	1.00 14.05	AAAA
MOTA	1734	CA	ASN	219	18.634	23.232	60.665	1.00 15.92	AAAA
MOTA	1735	CB	ASN	219	19.732	22.738	59.750	1.00 22.73	AAAA
MOTA	1736	CG		219	20.861	23.232	59.802	1.00 17.90	AAAA
ATOM	1737		ASN	219	19.398	21.789	58.868	1.00 16.62	AAAA
ATOM	1738	ND2		219	17.990	24.256	62.821	1.00 21.98	AAAA
ATOM	1739	C	ASN	219	17.075	23.569	63.262	1.00 18.65	AAAA
MOTA	1740	0	ASN	220	18.025	25.580	62.952	1.00 16.82	AAAA
ATOM	1741	N	ILE	220	16.951	26.298	63.640	1.00 13.22	AAAA
MOTA	1742	CA	ILE	220	17.522	27.115	64.823	1.00 15.70	AAAA
ATOM	1743	CB	ILE	220	16.411	27.912	65.479	1.00 15.18	AAAA
MOTA	1744		ILE	220	18.246	26.193	65.823	1.00 19.11	AAAA
MOTA	1745		ILE	220	17.350	25.259	66.632	1.00 22.75	AAAA .
ATOM -	1746		ILE	220	16.363	27.246	62.573	1.00 18.80	AAAA
MOTA	1747	C	ILE	220	16.810	28.386	62.419	1.00 15.52	AAAA
ATOM	1748	0	PRO	221	15.341	26.790	61.826	1.00 16.72	AAAA
MOTA	1749	И	PRO	221 .	14.612	25.518	61.906	1.00 18.83	AAAA
MOTA	1750	CD	PRO	221	14.739	27.628	60.785	1.00 19.83	AAAA
MOTA	1751	CA CB	PRO	221	13.930	26.615	59.948	1.00 20.76	AAAA
MOTA	1752	CG	PRO	221	14.409	25.241	60.462	1.00 28.73	AAAA
MOTA	1753	C	PRO	221	13.849	28.664	61.444	1.00 21.26	AAAA
MOTA	1754	ò	PRO	221	13.061	28.318	62.314	1.00 22.45	AAAA
ATOM	1755 1756	N	LEU	222	13.977	29.926	61.028	1.00 19.70	AAAA
ATOM	1757	CA	LEU	222	13.209	31.018	61.612	1.00 21.62	AAAA
ATOM	1758	CB	LEU	222	14.163	31.972	62.319	1.00 16.46	AAAA
MOTA	1759	CG	LEU	222	14.868	31.232	63.466	1.00 18.65	AAAA
MOTA	1760		LEU	222	16.026	32.072	64.014	1.00.21.32	AAAA
ATOM	1761		LEU	222	13.857	30.925	64.555	1.00 19.98	AAAA
ATOM	1762	C	LEU	222	12.350	31.763	60.590	1.00 19.68	AAAA
MOTA	1763	ō	LEU	222	12.687	31.830	59.412	1.00 18.07	AAAA
MOTA	1764	N	PRO	223	11.220	32.329	61.042	1.00 19.37	AAAA
MOTA	1765	CD	PRO	223	10.723	32.249	62.431	1.00 17.38	AAAA
MOTA	1766	CA	PRO	223	10.264	33.065	60.203	1.00 19.59	AAAA
MOTA	1767	CB	PRO	223	9.006	33.083	61.074	1.00 20.08	AAAA
MOTA	1768	CG	PRO	223	9.608	33.304	62.441	1.00 21.96	AAAA
MOTA		C	PRO	223	10.606	34.458	59.723	1.00 23.15	AAAA
ATOM	1769		PRO	223	11.525	35.101	60.214	1.00 15.81	AAAA
ATOM	1770	0	LYS	224	9.830	34.912	58.745	1.00 16.41	AAAA
ATOM	1771	N	LYS	224	9.975	36.254	58.200	1.00 16.11	AAAA
ATOM	1772	CA	LYS	224	9.002	36.446	57.039	1.00 20.34	AAAA
MOTA	1773	CB	LYS	224	9.163	35.441	55.900	1.00 19.33	AAAA
MOTA	1774	CG	LYS	224	8.109	35.687	54.807	1.00 25.49	AAAA
MOTA	1775	CD		224	8.209	34.624	53.704	1.00 24.14	AAAA
MOTA	1776	CE	LYS	224	7.207	34.843	52.618	1.00 34.08	AAAA
MOTA	1777	NZ	LYS	224	9.638	37.289	59.284	1.00 15.77	AAAA
MOTA	1778	C	LYS	224	8.819	37.032	60.186	1.00 21.07	AAAA
MOTA	1779	0	LYS GLY	225	10.239	38.469	59.171	1.00 20.77	AAAA
atom	1780	N	GLY	225	9.974	39.527	60.129	1.00 20.30	. AAAA
MOTA	1781	CA		225	10.556	39.286		1.00 20.63	AAAA
ATOM.	1782	С	GLY	ل مه ند	±0.330	,		_	_

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ATOM	1783	0	GLY	225	10.128	39.912	62.468	1.00 20.66	AAAA
ATOM	1784	N	LEU	226	11.540	38.395		1.00 20.37	AAAA
ATOM	1785				12.154	38.063	62.893	1.00 18.71	
									AAAA
ATOM	1786				13.354	37.145	62.670	1.00 13.63	AAAA
ATOM	1787				13.836	36.443	63.939	1.00 18.44	AAAA .
ATOM -	1788	CD	1 LEU	226	12.834	35.329	64.243	1,00 18.09	AAAA
MOTA	1789	CD	2 LEU	226	15.232	35.844	63.741	1.00 17.96	AAAA
MOTA	1790		LEU		12.649	39.309	63.642	1.00 19.84	ÀAAA
	1791		LEU		13.320	40.151			
ATOM							63.052	1.00 18.13	AAAA
ATOM	1792	N	ASN		12.336	39.421	64.932	1.00 23.30	AAAA
ATOM	1793	CA	ASN	227	12.815	40.571	65.692	1.00 20.88	AAAA
MOTA	1794	CB	ASN	.227	11.682	41.261	66.485	1.00 21.73	AAAA
MOTA	1795	CG	ASN	227	11.061	40.368	67.546	- 1.00 20.47	AAAA
ATOM	1796		1 ASN		11.762	39.736	68.341	1.00 23.80	
	_		2 ASN		9.729	40.340	67.581		AAAA
MOTA	1797							1.00 21.08	AAAA
MOTA	1798	С	ASN		13.950	40.152	66.612	1.00 25.24	- AAAA
MOTA	1799	0	ASN		14.282	38.965	66.702	1.00 18.54	AAAA
MOTA	1800	N	ASP	228	14.547	41.124	67.296	1.00 19.41	AAAA
ATOM	1801	CA	ASP	228	15.682	40.844	68.169	1.00 22.15	AAAA
ATOM	1802	CB	ASP		16.208	42.141	68.802	1.00 16.82	AAAA
ATOM	1803	CG	ASP		16.852	43.060	67.796	1.00 30.68	
				228	17.182	42.576			. AAAA
ATOM	1804		LASP				66.690	1.00 23.87	AAAA
ATOM	1805		2 ASP	228	17.053	44.256	68.123	1.00 25.02	AAAA
ATOM	1806	С	ASP	228	15.440	39.835	69.265	1.00 18.83	AAAA
MOTA	1807	0	ASP	228	16.298	39.002	69.536	1.00 16.28	AAAA
MOTA	1808	N	ASN	229	14.291	39.930	69.928	1.00 20.73	AAAA
MOTA	1809	CA	ASN	229	13.975	39.015	71.007	1.00 21.75	AAAA
ATOM	1810	CB	ASN	229	12.706	39.483	71.712	1.00 19.46	AAAA
ATOM	1811	CG	ASN	229	12.943	40.738	72.516	1.00 27.14	
									AAAA
ATOM	1812		ASN	229	13.588	40.691	73.556	1.00 33.03	AAAA
ATOM	1813		ASN	229	12.464	41.874	72.019	1.00 21.35	AAAA
ATOM	1814	С	ASN	229	13.833	37.596	70.503	1.00 18.47	AAAA
ATOM	1815	0	ASN	229	14.284	36.644	. 71.151	1.00 22.47	AAAA
ATOM	1816	N	GLU	230	13.252	37.454	69.319	1.00 17.79	AAAA
ATOM	1817	CA	GLU	230	13.081	36.125	68.748	1.00 21.18	AAAA
ATOM	1818	CB	GLU	. 230	12.152	36.193	67.536	1.00 20.54	AAAA
ATOM	1819	CG	GLU	230	10.765	36.714	67.890	1.00 28.98	
	1820	CD	GLU	230					AAAA
ATOM					9.870	36.816	66.677	1.00 24.35	AAAA
ATOM	1821		GLU	230	10.360	37.296	65.638	1.00 22.00	AAAA
MOTA	1822		GLU	230	8.683	36.443	66.772	1.00 24.99	AAAA
MOTA	1823	С	GLU	230	14.422	35.507	68.361	1.00 16.89	AAAA
MOTA	1824	0	GLU	230	14.663	34.326	68.603	1.00 19.45	AAAA
ATOM	1825	N	PHE	231	15.305	36.305	67.772	1.00 15.68	AAAA
ATOM	1826	CA	PHE	231	16.616	35.788	67.389	1.00 15.78	AAAA
ATOM	1827	CB	PHE	231	17.420	36.863	66.649	1.00 13.22	AAAA
	1828	CG	PHE	231	18.719	36.361	66.069	1 70.20.63	
MOTA									AAAA
MOTA	1829		PHE	231	18.723	35.445	65.016	1 00 18.42	AAAA
MOTA	1830		PHE	231	19.936	36.804	66.568	1 70 21.10	AAAA
MOTA	1831	CE1	PHE	231	19.918	34.983	64.471	1.00 17.67	AAAA
MOTA	1832	CE2	PHE	231	21.144	36.346	66.029	1.00 28.29	AAAA
MOTA	1833	CZ	PHE	231	21.130	35.431	64.976	1.00 27.85	AAAA
ATOM	1834	c	PHE	231	17.385	35.332	68.636	1.00 18.54	AAAA
MOTA	1835	0	PHE	231	17.869	34.201	68.702	1.00 18.86	AAAA
MOTA	1836	N	LEU	232	17.495	36.204	69,636	1.00 19.07	AAAA
ATOM	1837	CA	LEU	232	18.239	35.850	70.848	1.00 17.39	AAAA
MOTA	1838	CB	LEU	232	18.415	37.078	71.737	1.00 24.53	AAAA
ATOM	1839	CG	LEU	232	19.214	38.202	71.061	1.00 16.64	AAAA
ATOM	1840	CD1		232	19.134	39.449	71.934	1.00 26.70	AAAA
					20.659		70.810	1.00 18.77	AAAA
MOTA	1841	CD2		232		37.806			
ATOM	1842	С	LEU	232	17.607	34.707	71.628	1.00 19.82	AAAA
ATOM	1843	0	LEU	232	18.309	33.904	72.217	1.00 21.80	AAAA
ATOM	1844	N	PHE	233	16.281	34.640	71.648	1.00 17.18	AAAA
ATOM	1845	CA	PHE	233	15.587	33.537	72.309	1.00 23.34	AAAA
ATOM	1846	CB	PHE	233	14.074	33.663	72:095	1.00 19.17	AAAA
ATOM	1847	CG	PHE	233	13.289	32.447	72.523	1.00 21.40	AAAA
•					12.863		73.841	1.00 29.62	AAAA
ATOM	1848	CD1	FRE	233	12.003	32.302	12.041	1.00 23.02	www

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ATOM	1849	CD2	PHE	233	12.942	31.473	71.596	1.00 19.92	AAAA
ATOM	1850		PHE	233	12.088	31.206	74.229	1.00 29.35	AAAA
ATOM	1851		PHE	233	12.168	30.363	71.966	1.00 25.37	AAAA
ATOM	1852	CZ	PHE	233	11.737	30.231	73.283	1.00 30.28	AAAA
ATOM	1853	С	PHE	233	16.041	32.234	71.660	1.00 23.12	AAAA
ATOM	1854	0	PHE	233	16.433	31.273	72.332	1.00 18.35	AAAA
MOTA	1855	N	ALA	234	15.961	32.208	70.332	1.00 17.26	AAAA
ATOM	1856	CA	ALA	234	16.332	31.026	69.562	1.00 17.67	AAAA
ATOM	1857	CB	ALA	234	16.085	31.297	68.046	1.00 19.08	AAAA
ATOM	1858	С	ALA	234	17.786	30.641	69.800	1.00 16.31	AAAA
MOTA	1859	0	ALA	234	18.127	29.461	69.926	1.00 16.75	AAAA
ATOM	1860	N	LEU	235	18.646	31.643	69.846	1.00 16.73	AAAA
ATOM	1861	CA	LEU	235	20.074	31.411	70.051	1.00 19.14	AAAA .
MOTA	1862	CB	LEU	235	20.823	32.742	69.956	1.00 21.72 1.00 36.73	AAAA
ATOM	1863	CG	LEU	235	22.226	32.790	69.345 70.105	1.00 30.73	AAAA AAAA
MOTA	1864		LEU	235	23.026	33.844 31.426	69.393	1.00 22.96	AAAA
MOTA	1865		LEU	235	22.917 20.354	30.776	71.421	1.00 18.71	AAAA
ATOM	1866	C	LEU	235	21.028	29.747	71.522	1.00 18.59	AAAA
MOTA	1867	0	LEU	235 236	19.831	31.390	72.479	1.00 25.43	AAAA
ATOM	1868	N	GLU	236	20.046	30.883	73.839	1.00 19.75	AAAA
ATOM	1869	CA CB	GLU GLU	236	19.335	31.777	74.860	1.00 23.18	AAAA
MOTA	1870	CG	GLU	236	19.725	33.229	74.777	1.00 38.53	AAAA
MOTA	1871 1872	CD	GLU	236	18:857	34.119	75.648	1.00 42.42	AAAA
MOTA MOTA	1873		GLU	236	17.617	34.171	75.428	1.00 45.43	AAAA
ATOM	1874		GLU	236	19.425	34.768	76.548	1.00 48.76	AAAA
ATOM	1875	C	GLU	236	19.541	29.452	74.011	1.00 25.85	AAAA
ATOM	1876	ō	GLU	236	20.222	28.603	74.597	1.00 21.36	AAAA
ATOM	1877	Ŋ	LYS	237	18.343	29.193	73.501	1.00 23.16	AAAA
ATOM	1878	CA	LYS	237	17.752	27.871	73.610	1.00 17.06	AAAA
ATOM	1879	CB	LYS	237	16.282	27.943	73.193	1.00 26.98	AAAA
MOTA	1880	CG	LYS	237	15.483	26.711	73.519	1.00 52.00	AAAA AAAA
MOTA	1881	CD	LYS	237	14.078	27.110	73.932 75.183	1.00 56.40 1.00 52.03	AAAA
MOTA	1882	CE	LYS	237	14.131	27.979 28.421	75.614	1.00 55.53	AAAA
ATOM	1883	NZ	LYS	237	12.782 18.502	26.827	72.785	1.00 18.46	AAAA
MOTA	1884	C	LYS	237 237	18.691	25.692	73.231	1.00 21.20	AAAA
ATOM	1885	и	LYS SER	238	18.932	27.187	71.578	1.00 21.28	AAAA
MOTA	1886 1887	CA	SER	238	19.649	26.208	70.776	1.00 16.47	AAAA
ATOM ATOM	1888	CB	SER	238	19.745	26.666	69.307	1.00 19.75	AAAA
ATOM	1889	OG	SER	238	20.475	27.858	69.160	1.00 22.52	AAAA
MOTA	1890	C	SER	238	21.039	25.923	71.361	1.00 18.79	AAAA
ATOM	1891	0	SER	238	21.521	24.788	71.312	1.00 20.60	AAAA
ATOM	1892	N	LEU	239	21.690	26.937	71.925	1.00 22.95	AAAA
ATOM	1893	CA	LEU	239	23.004	26.701	72.513	1.00 20.98	AAAA
MOTA	1894	CB.	LEU	239	23.652	28.008	72.986	1.00 18.39	AAAA
ATOM	1895	CG	LEU	239	23.985	29.072	71.933	1.00 20.02 1.00 27.02	AAAA AAAA
ATOM	1896		LEU	239	24.538	30.311	72.636	1.00 27.02	AAAA
MOTA	1897	CD2		239	25.010	28.556	70.933 73.680	1.00 25.16	AAAA
MOTA	1898	c	LEU	239	22.882	25.735 24.929	73.920	1.00 20.70	AAAA
ATOM	1899	0	LEU	239	23.780 21.768	25.800	74.398	1.00 24.93	AAAA
MOTA	1900	N	GLU	240	21.700	24.912	75.536	1.00 25.72	AAAA
ATOM	1901	CA	GLU GLU	240 240	20.331	25.356	76.337	1.00 29.10	AAAA
ATOM	1902	CB CG	GLU	240	20.042	24.531	77.581	1.00 49.56	AAAA
ATOM	1903 1904	CD	GLU	240	19.053	25.212	78.515	1.00 63.15	AAAA
ATOM	1905	OE1		240	17.935	25.550	78.067	1.00 69.26	AAAA
ATOM	1905		GLU	240	19.400	25.410	79.703	1.00 66.68	AAAA
ATOM ATOM	1907	C	GLU	240	21.440	23.469	75.046	1.00 23.44	AAAA
ATOM	1908	ō	GLU	240	21.951	22.535	75.674	1.00 23.10	AAAA
MOTA	1909	N	ILE	241	20.771	23.294	73.913	1.00 19.52	AAAA
ATOM	1910	CA	ILE	241	20.598	21.978	73.321	1.00 24.06	AAAA
ATOM	1911	CB	ILE	241	19.705	22.039	72.052	1.00 23.80	AAAA
ATOM	1912	CG2	ILE	241	19.718	20.678	71.323	1.00 24.94	AAAA
ATOM	1913	CG1	ILE	241	18.281	22.433	72.439	1.00 28.60	AAAA
ATOM	1914	CD1	ILE	241	17.336	22.600	71.257	1.00 27.04	AAAA

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ATOM			ILI		21.95	7 21.404	72.941	1.00 25.48	AAAA
ATOM			IL		22.24		-	1.00 19.43	AAAA
ATOM	191		VAI		22.799				. AAAA
ATOM	191				24.116				
MOTA MOTA	1919 1920		3 VAI 31 VAI		24.853				
ATOM	192		31 VAL 32 VAL		26.273				· -
ATOM	192		VAL		24.093 24.962				
ATOM	1923		VAL		25.566				
ATOM	1924		LYS		24.989				
ATOM	1925				25.775			1.00 26.06 1.00 32.57	
ATOM	1926				25.599			1.00 32.57	
ATOM	1927				26.386		77.568	1.00 43.21	
MOTA	1928	CD	LYS	243	26.022		78.653	1.00 53.10	
MOTA	1929			243	26.407	25.607	78.287	1.00 50.30	
MOTA	1930				26.045	26.548	79.389	1.00 59.15	
MOTA	1931		LYS		25.433		76.046	1.00 30.38	
MOTA	1932		LYS		26.321		76.578	1.00 35.44	AAAA
ATOM ATOM	1933 1934		GLU		24.161		76.076	1.00 28.12	
ATOM	1935			244 244	23.798		76.798	1.00 37.54	AAAA
ATOM	1936			244	22.288 21.735		77.048	1.00 35.34	AAAA
ATOM	1937			244	20.281	20.275	77.816 78.230	1.00 55.88 1.00 57.89	AAAA
ATOM	1938		1 GLU	244	19.673	21.246	78.738	1.00 57.89	AAAA AAAA
ATOM	1939			244	19.753	19.152	78.062	1.00 57.73	AAAA
ATOM	1940	С	GLU	244	24.231	18.034	76.102	1.00 38.17	AAAA
MOTA	1941	0	GLU	244	24.294	16.978	76.727	1.00 38.46	AAAA
ATOM	1942	N	VAL	245	24.541	18.124	74.817	1.00 30.29	AAAA
MOTA	1943	CA	VAL	245	24.933	16.958	74.042	1.00 29.17	AAAA
MOTA MOTA	1944 1945	CB	VAL L VAL	245 245	23.984	16.778	72.833	1.00 46.68	AAAA
MOTA	1946	CG2		245	24.462 22.581	15.641 16.488	71.942	1.00 53.09	AAAA
ATOM	1947	c	VAL	245	26.364	16.982	73.327 73.508	1.00 54.19 1.00 34.90	AAAA
MOTA	1948	Ó	VAL	245	26.915	15.939	73.164	1.00 34.73	AAAA AAAA
MOTA	1949	N	PHE	246	26.980	18.156	73.465	1.00 29.22	AAAA
ATOM	1950	CA	PHE	246	28.324	18.256	72.897	1.00 29.17	AAAA
ATOM	1951	CB	PHE	246	28.178	18.800	71.464	1.00 30.42	AAAA
MOTA	1952	CG	PHE	246	29.384	18.588	70.585	1.00 25.62	AAAA
ATOM ATOM	1953 1954		PHE	246 246	29.695	17.326	70.097	1.00 28.89	AAAA
ATOM	1955		PHE	246	30.167 30.771	19.668 17.138	70.196 69.222	1.00 25.17	AAAA
ATOM	1956	CE2		246	31.248	19.495	69.322	1.00 23.43 1.00 22.40	AAAA
MOTA	1957	CZ	PHE	246	31.549	18.236	68.835	1.00 22.40	AAAA AAAA
MOTA	1958	C	PHE	246	29.233	19.176	73.712	1.00 23.38	AAAA
MOTA	1959	0	PHE	246	28.867	20.312	74.002	1.00 29.15	AAAA
ATOM	1960	N	GLU	247	30.410	18.682	74.094	1.00 29.73	AAAA
ATOM	1961	CA	GLU	247	31.395	19.481	74.841	1.00 28.10	AAAA
ATOM	1962	CB	GLU	247	31.912	18.726	76.074	1.00 35.75	AAAA
atom atom	1963 1964	CG CD	GLU GLU	247 247	30.972	18.707	77.286	1.00 60.78	AAAA
ATOM	1965		GLU	247	29.700 28.913	17.892 18.220	77.077 76.165	1.00 70.07	AAAA
ATOM	1966		GLU	247	29.481	16.920	77.835	1.00 79.95 1.00 76.80	AAAA
ATOM	1967	C	GLU	247	32.554	19.741	73.876	1.00 28.90	AAAA AAAA
ATOM	1968	0	GLU	247	33.490	18.946	73.778	1.00 23.67	AAAA
ATOM	1969	N	PRO	248	32.531	20.891	73.181	1.00 25.02	AAAA
MOTA	1970	CD	PRO	248	31.574	22.003	73.310	1.00 27.23	AAAA
ATOM	1971	CA	PRO	248	33.566	21.249	72.209	1.00 28.06	AAAA
MOTA	1972	CB	PRO	248	33.050	22.575	71.639	1.00 28.11	AAAA
ATOM	1973	CG	PRO	248	31.551	22.512	71.897	1.00 34.57	AAAA
ATOM	1974 1975	C	PRO	248	34.968	21.416	72.770	1.00 23.87	AAAA
ATOM	1975	N O	PRO GLU	248 249	35.132 35.965	21.897	73.887	1.00 24.05	AAAA
atom Atom	1977	CA	GLU	249	37.366	21.013 21.195	71.983 72.355	1.00 24.34	AAAA
	1978		GLU	249	38.275	20.166	71.679	1.00 25.98 1.00 22.07	AAAA AAAA
ATOM	1979		GLU	249	38.046	18.726	72.116	1.00 22.07	AAAA
MOTA	1980		GLU	249	39.005	17.767	71.445	1.00 29.15	AAAA
				٠			•		

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Figure 16-31

ATOM	1981	OE1	GLU	249	39.071	17.770	70.199	1.00 27.62	AAAA
	1982		GĻŪ	249	39.694	17.004	72.161	1.00 26.19	AAAA
MOTA						22.561	71.786	1.00 26.04	
MOTA	1983	С	GLU	249	37.692				AAAA
MOTA	1984	0	GLU	249	38.582	23.271	72.262	1.00 26.39	AAAA
ATOM	1985	N	VAL	250	36.953	22.921	70.744	1.00 23.83	AAAA
MOTA	1986	CA	VAL	250	37.151	24.197	70.086	1.00 19.67	AAAA
ATOM	1987	CB	VAL	250	38.438	24.178	69.210	1.00 20.88	AAAA
			-	250	38.348	23.117	68.128	1.00 18.18	AAAA
MOTA	1988		VAL						
MOTA	1989	CG2	VAL	.250	38.647	25.530	68.591	1.00 16.71	AAAA
ATOM	1990	С	VAL	250	35.946	24.483	69.207	1.00 20.78	AAAA
ATOM	1991	0	VAL	250	. 35.299	23.556	68.746	1.00 19.60	AAAA
ATOM	1992	N	TYR	251	35.633	25.757	69.000	1.00 18.75	AAAA
ATOM	1993	CA	TYR	251	34.497	26.109	68.153	1.00 22.44	ÄAAA
	1994	CB	TYR	251	33.261	26.437	69.022	1.00 16.57	AAAA
ATOM							69.575	1.00 22.36	
MOTA	1995	CG	TYR	251	33.207	27.856			AAAA
MOTA	1996		TYR .		32.654	28.896	68.823	1.00 18.12	AAAA
MOTA	1997	CEl	TYR	251	32.612	30.185	69.308	1.00 20.40	AAAA
ATOM	1998	CD2	TYR	251	33.715	28.160	70.842	1.00 20.04	AAAA
ATOM	1999	CE2	TYR	251	33.676	29.475	71.349	1.00 16.60	AAAA
ATOM	2000	CZ	TYR	251	33.128	30.473	70.573	1.00 14.68	AAAA
		ОН	TYR	251	33.100	31.780	71.011	1.00 21.79	AAAA
ATOM	2001					27.294	67.236	1.00 20.28	AAAA
MOTA	2002	C	TYR	251	34.811				
MOTA	2003	0	TYR	251	35.695	28.107	67.525	1.00 19.91	AAAA
ATOM	2004	N	LEU	252	34.097	27.360	66.109	1.00 17.90	AAAA
MOTA	2005	CA	LEU	252	34.216	28.466	65.161	1.00 18.58	AAAA
ATOM	2006	CB	LEU	252	34.679	28.001	63.767	1.00 17.55	AAAA
ATOM	2007	CG	LEU	252	36.028	27.290	63.718	1.00 23.36	AAAA
ATOM	2008		LEU	252	35.819	25.820	64.017	1.00 27.78	AAAA
					36.631	27.440	62.331	1.00 27.29	AAAA
ATOM	2009		LEU	252				1.00 27.29	AAAA
MOTA	2010	С	LEU	252	32.816	29.049	65.052		
ATOM	2011	0	LEU		31.819	28.320	65.120	1.00 18.82	AAAA
ATOM	2012	N	LEU	253	32.756	30.360	64.891	1.00 16.80	AAAA
MOTA	2013	CA	LEU	253	31.498	31.105	64.817	1.00 17.50	AAAA
ATOM	2014	CB	LEU	253	31.379	31. 9 87	66.073	1.00 15.49	AAAA
ATOM	2015	ĊG	LEU	253	30.326	33.085	66.165	1.00 17.75	AAAA
ATOM	2016		LEU	253	28.946	32.438	66.172	1.00 20.85	AAAA
	2017		LEU	253	30.536	33.897	67.464	1.00 19.05	AAAA
ATOM					31.516	31.985	63.580	1.00 20.22	AAAA
ATOM	2018	C	LEU	253					AAAA
MOTA	2019	0	LEU	253	32.474	32.727	63.371	1.00 18.14	
ATOM	2020	N	GLN	254	30.466	31.913	62.765	1.00 16.50	AAAA
ATOM	2021	CA	GLN	254	30.411	32.730	61.556	1.00 16.48	AAAA
ATOM	2022	CB	GLN	254	30.085	31.863	60.312	1.00 25.58	AAAA
ATOM	2023	CG	GLN	254	28.647	31.798	59.871	1.00 36.40	AAAA
ATOM	2024	CD	GLN	254	28.337	32.728	58.701	1.00 33.18	AAAA
	2025		GLN	254	28.744	32.487	57.546	1.00 21.05	AAAA
ATOM	2026		GLN	254	27.613	33.799	58.992	1.00 22.85	AAAA
ATOM					29.384	33.816	61.832	1.00 16.12	AAAA
ATOM	2027	C	GLN	254					AAAA
MOTA	20285	0	GLN	254	28.282	33.577	62.364	1.00 13.97	
ATOM	2029	N	LEU	255	29.768	35.032	61.468	1.00 14.42	AAAA
ATOM	2030	CA	LEU	255	28.988	36.215	61.763	1.00 17.99	AAAA
ATOM	2031	CB	LEU	255	29.834	37.070	62.719	1.00 20.68	AAAA
ATOM	2032	CG	LEU	255	30.240	36.283	63.964	1.00 22.90	AAAA
	2033		LEU	255	31.446	36.906	64.635	1.00 29.36	AAAA
MOTA				255	29.042	36.214		1.00 14.80	AAAA
ATOM	2034		LEU					1.00 19.32	AAAA
MOTA	2035	Ç	LEU	255	28.541	37.060	60.594		
ATOM	2036	0	LEU	255	28.838	38.260	60.561	1.00 21.23	AAAA
ATOM	2037	N	GLY	256	27.827	36.467	59.639	1.00 17.21	AAAA
MOTA	2038	CA	GLY	256	27.347	37.259	58.516	1.00 15.64	AAAA
ATOM	2039	C	GLY	256	26.413	38.348	59.028	1.00 17.31	AAAA
ATOM	2040	ō	GLY	256	25.717	38.150	60.027	1.00 15.62	AAAA
	2041	N	THR.	257	26.389	39.494	58.348	1.00 19.72	AAAA
ATOM				257	25.536	40.598	58.776	1.00 19.88	AAAA
ATOM	2042	CA	THR				58.589	1.00 14.02	AAAA
MOTA	2043	CB	THR	257	26.242	41.973			
ATOM	2044	OG1		257	26.538	42.187	57.206	1.00 17.58	AAAA
MOTA	2045	CG2	THR	257	27.543	42.009	59.392	1.00 19.67	AAAA
ATOM	2046	С	THR	257	24.199	40.634	58.053	1.00 20.58	AAAA

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ATOM			THE		23.40	03 41.545	58.266	1.0	0 14.59	AAAA
MOTA MOTA			ASF		23.92			1.0	0 16.56	AAAA
ATOM					22.65 22.60			_		AAAA
ATOM					23.03				0 18.38 0 25.85	AAAA
ATOM	- 205	2 OD	1 ASP		23.22				0 22.32	AAAA AAAA
MOTA	205		2 ASP		23.18	37 36.370		-	0 18.12	AAAA
MOTA			ASP		21.39			1.0	0 21.25	ÄAAA
ATOM ATOM	205 205		ASP PRO		20.30				0 22.52	AAAA
ATOM	205				21.51 22.61				0 18.17 0 25.88	AAAA
ATOM	205		PRO	259	20.28			1.0	0 23.88	AAAA AAAA
MOTA	205		PRO	259	20.71			- 1.0	0 21.18	AAAA
ATOM	206	-	PRO	259	22.17		60.846	1.0	0 36.11	AAAA
ATOM ATOM	206 206		PRO PRO	259 259	19.70 18.57				0 20.88	AAAA
ATOM	206		LEU	260	20.47				0 19.25 0 18.75	-AAAA
ATOM	206		LEU	260	20.02		59.875			AAAA AAAA
ATOM	206		LEU	260	21.20	2 43.935			0 20.35	AAAA
MOTA	2066		LEU.		22.40		60.682	1.0	21.82	AAAA
MOTA MOTA	2067 2068		LEU	260 260	23.60 22.03		60.253		18.57	AAAA
ATOM	2069		LEU	260	18.87		62.123 59.014	1.00		AAAA
MOTA	2070		LEU	260	18.74		57.826		21.69	AAAA AAAA
ATOM	2071		LEU	261	18.04	9 44.300	59.634		19.54	AAAA
ATOM ATOM	2072 2073		LEU LEU	261	16.90		58.965		17.34	AAAA
ATOM	2074		LEU	261 261	16.28 15.20		59.892 59.300		19.96	AAAA
ATOM	2075		LEU	261	14.08		58.732		29.99 33.66	AAAA AAAA
MOTA	2076		LEU	261	14.682		60.376		44.71	AAAA
ATOM	2077		LEU	261	17.262		57.620	1.00	18.11	AAAA
ATOM ATOM	2078 2079		LEU GLU	261 262	16.539 18.391		56.634	1.00		AAAA
ATOM	2080		GLU	262 .			57.566 56.338	1.00	22.68	AAAA AAAA
ATOM	2081		GLU	262	19.875		56.641		22.01	AAAA
MOTA	2082		GLU	262	19.365		57.443	1.00	22.94	AAAA
ATOM ATOM	2083 2084	CD OE1	GLU GLU	262 262	19.434 19.668		58.927		23.11	AAAA
ATOM	2085		GLU	262	19.238		59.357 59.667	1.00	24.58	AAAA
ATOM	2086	С	GLU	262	19.281		55.197		25.65	AAAA AAAA
MOTA	2087	0	GLU	262	19.446		54.070		25.49	AAAA
ATOM ATOM	2088 2089	N CA	ASP ASP	263 263	19.501		55.467		22.45	AAAA
ATOM	2090	CB	ASP	263	19.959 20.981		54.418 54.988		15.93 18.99	AAAA
ATOM	2091	CG	ASP	263	21.706		53.907		22.21	AAAA AAAA
MOTA	2092	OD1		263	22.876	41.730	54.139		23.19	AAAA
ATOM ATOM	2093 2094	OD2 C	ASP ASP	263	21.112		52.838	1.00		AAAA ·
ATOM	2095		ASP	263 263	18.733 18.012		53.837 54.519		22.32	AAAA
ATOM	2096		TYR	264	18.500		52.564		18.50 25.21	AAAA AAAA
MOTA	2097	CA	TYR	264	17.339	42.936	51.865		29.92	AAAA
ATOM	2098		TYR	264	17.077		50.596	1.00	38.48	AAAA
MOTA MOTA	2099 2100	CG CD1	TYR	264 264	17.910		49.379		54.09	AAAA
MOTA	2101	CE1		264	17.677 18.420	42.249 41.930	48.660 47.526		69.38	AAAA
ATOM	2102	CD2	TYR	264	18.915	44.286	48.928		68.71 66.09	AAAA AAAA
MOTA	2103	CE2		264	19.670	43.975	47.788		74.50	AAAA
ATOM	2104		TYR	264	19.415	42.794	47.094	1.00	72.57	AAAA
MOTA MOTA	2105 2106		TYR TYR	264 264	20.154 17.445	42.472	45.975		71.96	AAAA
ATOM	2107		TYR	264	16.448	41.461 40.839	51.532 51.190		29.55 30.11	AAAA
MOTA	2108		LEU	265	18.639	40.891	51.629		24.45	AAAA AAAA
MOTA	2109		EU	265	18.753	39.476	51.337		25.36	AAAA
MOTA	2110		EU	265	20.186	39.089	50.969		29.81	AAAA
MOTA MOTA	2111	CG I	EU	265 265	20.509 21.847	39.510	49.531		34.43	AAAA
11011	تكلين	ent i) III		21.04/	38.930	49.100	1.00	44.38	AAAA

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Figu	te 10-33	
- 0	10 603 1 00 46.72	AAAA
sier 1	9 422 38,990 30.447 1.00 22.33	AAAA
NOW 2113 CD2 LEU 205	8.209 38.585 52.348 1.00 23.48	AAAA
2114 C LEU 205	18.279 37.364 52.508 1.00 17.50	AAAA AAAA
2115 O LEU 203	17.677 39.194 54.569 1.00 19.69	•
MOM 2116 N SER 200	17.055 38.398 55.845 1.00 20.73	AAAA AAAA
2117 CA SER 200	17 912 38.314 33.604 1 00 22.81	AAAA
2118 CB SER 200	17.696 39.442 54 950 1.00 19.75	AAAA
100 2119 OG SER 200	15.739 39.046 54.840 1.00 23.66	AAAA
2120 C SER 200	15.572 40.265 55.402 1.00 18.40	AAAA
2121 0 SER 200	14.799 38.223 55 851 1.00 20.64	AAAA
AMOM 2122 N 510 267	13.527 38.755 55 513 1.00 20.96	AAAA
атом 2123 СА 515 267	10 39/ 3/*** 608 1 00 20.00	AAAA
ATOM 2124 CB LIS 267	12.269	AAAA
ATOM 2125 CG 247	17 (193 22*** 684 1 (1) 30.72	AAAA
ATTOM 2126 CD 525 367	11.985 30 793 50.991 1.00 33.42	AAAA
ATOM 2127 CE 515 267	11.954 39 987 57.365 1.00 20.03	AAAA
ATOM 2128 NZ 257	13.601 39.192 58.017 1.00 25.30	AAAA
ATOM 2129 C 175 267	12.584 39.437 57.915 1.00 18.50	AAAA
ATOM 2130 0 DUE 268	14.814 30.182 59.345 1.00 10.30	AAAA
ATOM 2131 N PUE 268	13.05 20 510 59.833 1.00 1.06	AAAA
ATOM 2132 THE 268	10.32 22 006 59.907 1.02 62	AAAA
ATOM 2133 CB PUE 268	16.252 36 290 60.415 1.00 10.13	AAAA
ATOM 2134 CD DUE 268	16 091 36.303 59.002 1 00 14.81	AAAA
ATOM CD2 PHE 200	17.331 34.904 60.381 1.00 17.45	AAAA
ATOM CEI PHE 200	15 027 34.900 59.849 1.00 16.01	AAAA
ATOM 2139 CE2 PHE 200	16 144 34.208 60.296 1.00 18.33	AAAA
ATOM 2130 CZ PHE 268	15 179 40.699 59.510 1 00 18.28	AAAA
ATOM 2140 C PHE 268	15 733 41.3/1 30.613 1 00 21.04	AAAA AAAA
ATOM 2141 O PHE 268	14 679 41.238 050 1 00 22.89	AAAA AAAA
ALON ASIN DO	14.763 42.673 60.940 1.00 20.55	AAAA
2143 CA ASN	13.365 43.298 50.686 1.00 26.13	AAAA
2144 CB ASN 200	12 551 43.0/1.	AAAA
ATOM 2145 CG ASN 269	13.060 43.192 50.860 1.00 28.26	AAAA
MOOM 2146 ODI ASN 269	11.200 42 159 1.00 13.00	AAAA
ATOM 214/ ND2 1.50	15.493 42.683 63.019 1.00 21.03	AAAA
MOM 2148	14.984 43.435 62.298 1.00 17.57	AAAA
NTOM 2149 0 750 270	16.695 42.642 63.521 1.00 10.05	AAAA
ATOM 2150 N 220 270	43 507 63.712 4.50 20 54	AAAA
NTOM 4131 370	17 045 40.058 63.631 1.00 14.19	AAAA
ATOM 2132 - 1511 270	10,070 39,174 64,132 1,00 19,05	AAAA
ATOM 2153 CD1 LEU 270	16 679 39.853 64.463 1 00 22.83	AAAA
ATOM 270	19 203 43.971 63.560 1 00 18.25	AAAA.
ATOM SIES C LEU 270	10 409 44.643 62.500 1 00 20.95	AAAA AAAA
ATOM 2157 O LEU 270	19 621 44.310 03.00 100 18.28	
2158 N SER 071	19.414 45.518 66.409 1.00 18.73	
2159 CA SER 271	18.985 46.130 67.512 1.00 22.28	
200 CB SER 271	19.347 45.327 55 224 1.00 19.96	
NTOM 2161 00 271	20.8/5 45.50 65 537 1.00 10.01	AAAA
ATOM 2162 C SER 271	21.122 45.004 65.020 1.00 10.2	AAAA
NTOM 2163 0 272	21.828 45.695 65.145 1.00 20.4	y
ATOM 2164 N 272 272	23.270 45.003 64.884 1.00 57.5	•
ATOM 2165 CA 150 272	24.176 47 378 63.483 1.00 34.6	7
ATOM 2166 CD 272 272	40 454 63.13	
ATOM 2167 CG 1151 272	46 504 62.373	
ATOM 2100 NID2 ASN 272	22 586 45.343	
ATOM 2103 C ASN 272	24 545 44.625 66.834 1.00 19.5	,
ATOM 2170 C ASN 272	22 231 45.930 0 22.	14
ATOM 2171 N VAL 273	22 053 45.698 30 755 1 00 26.	91
ATOM 21/2 C3 VAL 273	22 345 46./65 23.022 1 00 39.	63
ATOM 2173 CB VAL 273	22 440 46.444 1 2 2 1 00 34.	13
ATOM 2174 CG1 VAL 273	22 034 48.115 69.331 1 00 22.	06
ATOM 2176 CG2 VAL 27	22 636 44.293	89 AAAA
ATOM 221	22 249 .43 .708 /0.21	•
ATO O VAL	, , ,	
ATOM 2178 0		
•		

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MOTA	217		ALA	274	21.601	43.747	68.713	1.00 21.79	AAAA
ATOM	218	0 C.	A ALA	274	21.207	42.383	69.035		AAAA
ATOM	218	1 C	B ALA	274	19.806				AAAA
ATOM	218	2 C	ALA	274	22.259				AAAA
ATOM	218	30	ALA	274	22.569				AAAA
ATOM	218	4 N	PHE	275	22.798				AAAA
ATOM	218	5 C	A PHE	275	23.828				AAAA
ATOM	218	6 CI	B PHE		24.220				AAAA
ATOM	218	7 C	PHE	275	25.363				AAAA
ATOM	218		ol PHE	_	25.209				
ATOM	2189		2 PHE		26.590				AAAA
ATOM	2190		1 PHE		26.266				AAAA
ATOM	219		2 PHE		27.654				AAAA
ATOM	2192				27.489				AAAA
ATOM	2193		PHE	_				1.00 25.06	AAAA
ATOM	2194		PHE		25.619			1.00 23.00	AAAA
ATOM	2195		LEU	276	25.366			1.00 17.49	AAAA
ATOM	2196			276	26.482			1.00 24.23	AAAA
ATOM	2197			276	26.736			1.00 20.44	AAAA
ATOM	2198			276	28.001	43.967		1.00 20.44	AAAA
ATOM	2199		1 LEU	276	27.948	45.447		1.00 29.65	AAAA
ATOM	2200		2 LEU	276	28.102	43.143	71.460		AAAA
ATOM	2201		LEU	276	26.180	41.278	70.262	1.00 32.41	AAAA
ATOM	2202		LEU	276	27.045	40.529	70.727	1.00 18.88	AAAA
ATOM	2203	-	LYS	277	24.968	41.374	70.727	1.00 17.99 1.00 19.67	AAAA
ATOM	2204			277	24.644	40.552	71.964	1.00 19.67	AAAA
ATOM	2205		LYS	277	23.265	40.888	72.532	1.00 21.33	AAAA
ATOM	2206		LYS	277	23.247	42.126	73.366	1.00 23.84	AAAA
ATOM	2207	CD	LYS	277	22.069	42.086	74.325	1.00 54.73	AAAA
MOTA	2208	CE	LYS	277	.22.172	40.884	75.254	1.00 58.85	AAAA
ATOM	2209	NZ	LYS	277	21.051	40.844	76.228	1.00 55.34	AAAA
ATOM	2210	С	LYS	277	24.695	39.068	71.660	1.00 22.12	AAAA AAAA
ATOM	2211	0	LYS	277	25.074	38.264	72.513	1.00 22.12	AAAA
ATOM	2212	N	ALA	278	24.311	38.700	70.441	1.00 20.23	AAAA
ATOM	.2213	CA	ALA	278	24.325	37.291	70.039	1.00 17.06	AAAA
ATOM	2214	CB	ALA	278	23.798	37.154	68.589	1.00 19.27	AAAA
ATOM	2215	С	ALA	278	25.760	36.767	70.127	1.00 16:94	AAAA
ATOM	2216	0	ALA	278	26.035	35.676	70.648	1.00 14.93	AAAA
ATOM	2217	N	PHE	279	26.679	37.564	69.606	1.00 18.88	AAAA
ATOM	2218	CA	PHE	279	28.099	37.231	69.626	1.00 21.01	AAAA
ATOM	2219	CB	PHE	279	28.880	38.392	68.998	1.00 16.79	AAAA
ATOM	2220	CG	PHE	279	30.370	38.264	69.120	1.00 20.23	AAAA
MOTA	2221		PHE	279	31.062	37.272	68.423	1.00 21.61	AAAA
MOTA	2222		PHE	279	31.088	39.159	69.905	1.00 23.24	AAAA
ATOM	2223		PHE	279	32.461	37.185	68.509	1.00 30.98	AAAA
ATOM	2224		PHE	279	32.480	39.081	69.995	1.00 24.82	AAAA
MOTA	2225	CZ	PHE	279	33.169	38.095	69.295	1.00 30.27	AAAA
ATOM	2226	C	PHE	279	28.576	36.995	71.067	1.00 25.48	AAAA
ATOM	2227	0	PHE	279	29.275	36.016	71.362	1.00 16.30	AAAA
MOTA	2228	N	ASN	280	28.194	37.898	71.962	1.00 22.30	AAAA
ATOM	2229	CA	ASN	280	28.599	37.777	73.352	1.00 24.49	AAAA
ATOM	2230	CB	ASN	280	28.391	39.109	74.080	1.00 27.17	AAAA
ATOM	2231	CG	ASN	280	29.344	40.183	73.578	1.00 20.88	AAAA
ATOM	2232		ASN	280	30.503	39.897	73.273	1.00 22.95	AAAA
ATOM	2233		ASN	280	28.875	41.421	73.522	1.00 27.85	AAAA
ATOM	2234	С	ASN	280	27.928	36.636	74.095	1.00 23.01	AAAA
ATOM	2235	0	ASN	280	28.510	36.062	75.016	1.00 21.91	AAAA
ATOM	2236	N	ILE	281	26.711	36.300	73.689	1.00 18.74	AAAA
ATOM	2237	CA	ILE	281	26.005	35.179	74.294	1.00 18.37	AAAA
MOTA	2238	СВ	ILE	281	24.566	35.067	73.758	1.00 19.31	AAAA
MOTA	2239	CG2	ILE	281	23.977	33.725	74.135	1.00 28.87	AAAA
ATOM	2240	CG1	ILE	281	23.710	36.206	74.308	1.00 23.51	AAAA
MOTA	2241	CD1	ILE	281	22.279	36.193	73.776	1.00 26.47	AAAA
ATOM ·	2242	С	ILE	281	26.743	33.876	73.965	1.00 18.54	AAAA
MOTA	2243	0	ILE	281	26.830	32.973	74.801	1.00 19.69	AAAA
ATOM	2244	N	VAL	282	27.258	33.765	72.744	1.00 17.72	AAAA
							-	•	

53/263 Figure 16-35

ATOM	2245	CA	VAL	282	27.976	32,553	72.352	1.00	14.89	AAAA
						32.565	70.852	1.00	10 50	
ATOM	2246	CB	VAL	282	28.359					AAAA
ATOM	2247	CG1	VAL	282	29.342	31.440	70.567	1.00	20.73	AAAA
					27.105	32.363	69.994	1.00	17 49	AAAA
ATOM	2248	CGZ	VAL	282						
MOTA	2249	С	VAL	282	29.241	32.433	73.198	1.00	21.79	AAAA
		0	VAL	282	29.568	31.360	73.715	1.00	25 80	AAAA
ATOM	2250									
MOTA	2251	N	ARG	283	29.935	33.549	73.361	1.00	19.14	AAAA
	2252.	CA	ARG	283	31.161	33.548	74.150	1.00	23 51	AAAA
ATOM										
MOTA	2253	CB	ARG	283	31.851	34.898	74.023	1.00	20.64	AAAA
ATOM	2254	CG	ARG	283	32.338	35.200	72.607	1.00	19.65	AAAA
MOTA	2255	$^{\rm CD}$	ARG	283	32.754	36.645	72.474	1.00		AAAA
ATOM	2256	NE	ARG	283	33.970	36.944	73.215	1.00	36.05	AAAA
					34.277	38.147	73.681	1.00		AAAA
ATOM	2257	cz	ARG	283						
ATOM	2258	NH1	ARG	283	33.448	39.169	73.488	1.00	35.23	AAAA
	2259	MILES	ARG	283	35.419	38.332	74.326	1.00	29.30	AAAA
ATOM										
MOTA	2260	С	ARG	283	30.911	33.219	75.622	1.00	25.44	AAAA
ATOM	2261	0	ARG	283	31.754	32.600	76.272	1.00	23.12	AAAA
ATOM	2262	N	GLU	284	29.765	33.632	76.151	1.00		AAAA
ATOM	2263	CA	GLU	284	29.462	33.338	77.553	1.00	31.77	AAAA
			GLU	284	28.243	34.115	78.033	1.00	30 06	AAAA
ATOM	2264	CB								
ATOM	2265	CG	GLU	284	28.399	35.605	77.957	1.00	50.56	AAAA
	2266	CD	GLU	284	27.137	36.320	78.365	1.00	63.75	AAAA
ATOM								1.00		
MOTA	2267	OEl	GLU	284	26.085	36.067	77.738			AAAA
ATOM	2268	OE2	GLU	284	27.198	37.133	79.309	1.00	72.01	AAAA
					29.181	31.862	77.733	1.00		AAAA
MOTA	2269	С	GLU	284						
ATOM	2270	0	GLU	284	29.410	31.310	78.803	1.00	33.08	AAAA
	2271	N	VAL	285	28.673	31.221	76.686	1.00	23.37	AAAA
ATOM										
ATOM	2272	CA	VAL	285	28.354	29.807	76.774	1.00		AAAA
ATOM	2273	CB	VAL	285	27.221	29.407	75.789	1.00	24.77	AAAA
					26.952	27.913	75.881	1.00	25 98	AAAA
MOTA	2274		VAL	285						
MOTA	2275	CG2	VAL	285	25.940	30.181	76.107	1.00	24.98	AAAA
MOTA	2276	С	VAL	285	29.567	28.942	76.479	1.00	31.41	AAAA
								1.00		AAAA
MOTA	2277 -	0	VAL	285	29.833	27.983	77.195			
ATOM	2278	N	PHE	286	30.316	29.276	. 75.431	1.00	27.27	AAAA
	2279	CA	PHE	286	31.463	28.457	75.086	1.00	22.47	AAAA
ATOM										
ATOM	2280	CB	PHE	286	31.289	27.904	73.667	1.00		AAAA
ATOM	2281	CG	PHE	286	30.168	26.918	73.536	1.00	25.71	AAAA
					28.971	27.274	72.917	1.00	22 88	AAAA
MOTA	2282		PHE	286						
ATOM	2283	CD2	PHE	286	30.294	25.631	74.069	1.00	24.49	AAAA
	2284	CE1	PHE	286	27.919	26.365	72.829	1.00	19.85	AAAA
ATOM .								1.00		AAAA
ATOM	2285	CEZ	PHE	286	29.246	24.714	73.987			
ATOM	2286	CZ	PHE	286	28.056	25.081	73.367	1.00	24.59	AAAA
		c	PHE	286	32.854	29.059	75.225	1.00	21.53	AAAA
ATOM	2287									
ATOM	2288	0	PHE	286	33.849	28.417	74.873	1.00		AAAA
ATOM	2289	N	GLY	287	32.937	30.272	75.754	1.00	23.76	AAAA
							75 901	1.00	24 17	AAAA
MOTA	2290	CA	GLY	287	34.237	30.896				
ATOM	2291	С	GLY	287	34.705	31.419	74.562	1.00	27.05	AAAA
	2292	ō	GLY	287	33.888	31.670	.73.667	1.00	18.06	AAAA
MOTA										
ATOM	2293	N	GLU	288	36.017	31.576	74.414	1.00		AAAA
MOTA	2294	CA	GLU	288	36.583	32.085	73.170	1.00	24.87	AAAA
							73.410	1.00	20 25	AAAA
MOTA	2295	CB	GLU	288	37.968	32.682				
ATOM	2296	CG	GLU	288	37.984	33.933	74.291	1.00	42.63	AAAA
	2297	CD	GLU	288	37.114	35.052	73.745	1.00	43.77	AAAA
ATOM										
ATOM	2298	OE1	GLU	288	37.235	35.380	72.544	1.00	30.82	AAAA
ATOM	2299		GLU	288	36.317	35.617	74.521	1.00	51.56	AAAA
					36.693	31.028	72.072	1.00		AAAA
MOTA	2300	С	GLU	288						
ATOM	2301	0	GLU	288	36.995	29.856	72.332	1.00	T8.10	AAAA
		N	GLY	289	36.447	31.468	70.843	1.00	26.12	AAAA
ATOM	2302									
ATOM	2303	CA	GLY	289	36.517	30.588	69.692	1.00		AAAA.
	2304	C	GLY	289	37.126	31.318	68.510	1.00	18.56	AAAA
MOTA							68.679	1.00		AAAA
MOTA	2305	0	GLY	289	37.669	32.404				
ATOM	2306	N	VAL	290	37.032	30.724	67.322	1.00	19.86	AAAA
			VAL	290	37.572	31.312	66.103	1.00	19.70	AAAA
MOTA	2307	CA						1.00		AAAA
ATOM	2308	CB	VAL	290	38.150	30.192	65.184			
	2309	CG1	VAL .	290	38.667	30.769	63.853	1.00	15.54	AAAA
MOTA			178 Y	290				1.00		AAAA
MOTA	2310	CG2	٧AL	450	39.296	29.483 °	03.340	1.00	~ 4.40	Les ALTA

54/263 Figure 16-36

ATOM			: VA	L 290	36.4	408 32.0	40 65.42	7 1 (0 20.90	3333
ATOM) VA	և 290	35.3	351 31.43	39 65.19		00 19.33	AAAA
MOTA	231	.3 N	TY!	R 291	36.5				0 15.37	AAAA
ATOM	231	4 C	A TY		35.5			4 1.0	10 15.37	AAAA
ATOM	231		B TY		35.4			4 1.0	0 16.79	AAAA
ATOM		_	G TY						0 16.42	AAAA
ATOM					35.3				0 18.60	AAAA
		_	D1 TY		36.3			9 1.0	0 21.77	AAAA
MOTA			E1 TYP		36.3		35 69.036	1.0	0 22.55	AAAA
ATOM			D2 TYF	₹ . 291	34.3	88 34.37			0 17.34	
ATOM	232	0 с	E2 TYF	291	34.3				0 20.24	AAAA
MOTA	232	1 C	Z TYF		- 35.3					AAAA
ATOM	232				35.3	30 34 34			0 25.85	AAAA
ATOM								-	0 25.57	, AAAA
		-			35.7				0 14.97	AAAA
ATOM	232				36.7		1 62.586	1.0	0 15.21	AAAA
ATOM	232				34.6	60 34.18	9 62.273		0 14.06	AAAA
MOTA	232		A LEU	292	34.6	74 34.39	2 60.824	1.0	0 15.03	AAAA
MOTA	232	7 CI	B LEU	292	34.4	61 33.04			0 13.66	
MOTA	2328	3 C	G LEU	292	35.3				0 19.04	AAAA
MOTA	2329	CI	Ol LEU		34.9					AAAA
ATOM	2330		D2 LEU		36.7				0 15.17	AAAA
ATOM	2331		LEU						0 19.18	AAAA
					33.5				0 16.62	AAAA
ATOM	2332		LEU		32.5			1.0	0 14.76	AAAA
MOTA	2333		GLY		33.7	24 35.93	2 59.216		0 19.62	AAAA
ATOM	2334		CLY	293	32.6	96 36.81			17.10	AAAA
MOTA	2335	C	GLY	293	.31.6	11 35.95			23.44	
MOTA	2336	C	GLY	293	31.40				23.60	AAAA
ATOM	2337	N	GLY	294	30.93					AAAA
MOTA	2338			294	29.87				24.96	AAAA
ATOM	2339		GLY	294	29.13				27.07	AAAA
ATOM	2340		GLY	294					28.41	AAAA
MOTA	2341	N			29.60				25.66	AAAA
			GLY.	295	27.97			1.00	20.33	AAAA
ATOM	2342	CA		295	27.16		54.085	1.00	20.14	AAAA
ATOM	2343	С	GLY	295	26.74	2 38.244	54.730	1.00	25.34	AAAA
ATOM	2344	.0	GLY	295	26.55	0 38.317	55.942		28.89	AAAA
ATOM	2345	N	GLY	296	26.61				28.52	AAAA
ATOM	2346	CA	GLY	296	26.23	0 40.598			23.21	
ATOM	2347	C	GLY	296	26.31	4 41.342				AAAA
ATOM	2348	0	GLY	296	27.35				26.34	AAAA
MOTA	2349	N	TYR	297	25.23				26.05	AAAA
ATOM	2350	CA	TYR	297					22.61	AAAA
ATOM	2351	СВ	TYR		25.22				22.58	AAAA
ATOM	2352			297	24.26			1.00	23.68	AAAA
		CG	TYR	297	24.50			1.00	25.14	AAAA
MOTA	2353		LTYR	297	23.98		51.568	1.00	28.31	AAAA
MOTA	2354		TYR	297	24.26		51.662	1.00	24.18	AAAA
MOTA	2355		TYR	297	25.30	7 39.725	49.577		29.74	AAAA
MOTA	2: 56	CE2	TYR	297	25.59	8 38.362			27.09	AAAA
ATOM	257	cz	TYR	297	25.08		50.696		28.68	AAAA
MOTA	2538	ОН	TYR	297	25.40	7 36.261			28.17	AAAA
ATOM	2359	С	TYR	297	24.91		51.320			AAAA
ATOM	2360	0	TYR	297	24.841				24.98	AAAA
ATOM	2361	N.	HIS	298	24.740	· · - ·	50.237		26.51	AAAA
ATOM	2362						52.491		23.80	AAAA
		CA	HIS	298	24.480		52.591	1.00	23.44	AAAA
ATOM	2363		HIS	298	23.325		53.536	1.00	23.37	AAAA
ATOM	2364	CG	HIS	298	22.956	47.945	53.551	1.00	32.94	AAAA
MOTA	2365	CD2	HIS	298	23.491	48.983	54.232		24.31	AAAA
ATOM	2366	ND1	HIS	298	22.011		52.707		38.29	AAAA
ATOM	2367	CE1	HIS	298	21.978		52.868		26.60	
ATOM	2368		HIS	298	22.867					AAAA
ATOM	2369	C	HIS	298	25.757		53.788		36.57	AAAA
ATOM	2370	ō	HIS				53.184		23.11	AAAA
ATOM				298	26.135		54.306		22.64	AAAA
	2371	N	PRO	299	26.430		52.445		21.07	AAAA
ATOM	2372	CD	PRO	299	26.078		51.117	1.00	27.42-	AAAA
ATOM	2373	CA	PRO	299	27.676	48.286	52.910		26.62	AAAA
ATOM ·	2374	CB	PRO	299	28.041	49.228	51.755		28.95	AAAA
ATOM	2375	CG	PRO	299	26.678	49.600	51.196		35.16	AAAA
MOTA	2376	С	PRO	299	27.644	48.991	54.262		25.75	
	-	-	-	·		-0.091	23.505	1.00	23.13	AAAA

SUBSTITUTE SHEET (RULE 26)

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				Fig	ure 16-3/					
				•			55.068	1.00 24	. 30	AAA AAA
				299		18.845	54.504	1 00 24	.48 4	AAA AAA
MOTA	2377	v	PRO	300	26.602	49.769	55.766	1 00 22	.94	AAAA
MOTA	2378		TYR	300	26.495	50.478	55.734	1.00 25	, 43	AAAA
MOTA	2379		TYR TYR	300		51.442 52.599	54.762	1.00 30	. 44	AAAA
ATOM	2380		TYR	300		52.634	53.746	1.00 26		AAAA
MOTA	2381	CD1		300	20.9-	53.676	52.819	1.00 29	.00	AAAA
MOTA	2382	CE1	TIN	300	20.0	53.640	54.827	1.00 31		AAAA
ATOM	2383	CD2	TIV.	300	24.490	54.677	53.916	1.00 35	0.85	AAAA
MOTA	2384	CDZ	TYR	300	24.501	54.689	52.913	1.00 3	3.44	AAAA
MOTA	23.85		TYR	300	25.448	55.700	51.990	1.00 3	3.41	AAAA
MOTA	2386		TYR	300	25.417	49.515	56.921	1.00 2	2.80	AAAA
MOTA	2387	OH	TYR	300	26.280	49.643	57:983	1.00 1	2 UB	AAAA
MOTA	2388	C	TYR	300 .	26.895	48.568	56.705	1.00 2	1 68	AAAA
MOTA	2389	И	ALA	301	25.374	47.589	57.719	1.00 2	a 52 -	AAAA
MOTA	2390	CA	ALA	301	25.009	46.687	57.198		3 49	AAAA
MOTA	2391		ALA	301	23.893	46.762	58.090		21 21	AAAA
MOTA.	2392		ALA	301	26.216	46.570	59.274		3 19	AAAA
MOTA	2393	_	ALA	301	26.507 26.904	46.275	57.074		20.66	AAAA
MOTA	2394		LEU	302	28.090	45.463	57.234		23.31	AAAA
MOTA	2395		LEU	302	28.602	45.057	7 55.047		36.66	AAAA
MOTA	2396	· ~~		. 302	29.932	44.33	5 55.01.		38.41	AAAA
MOTA	2397 2398			302	29.979	43.84	9 54.1/		28.52	AAAA
ATOM		CE	1 LEU	302	31.104	45.25	5 55.87	1 00	22.08	AAAA
MOTA			2 LEU	302	29.165	46.20	4 58.01	0 1 00	20.43	AAAA
MOTA		_	LEU	302	29.653	45./1	3 59.02 1 57.54	0 1 00	19.58	AAAA AAAA
MOTA			LEU		29.517	47.40		7 1 00	19.77	AAAA
MOTA	240		ALA	7 7 7	30.567	, 48.T		2 1 00	21.69	AAAA
ATOTA	40			202	30.816	5 49.40		1 00	19.19	AAAA
IOTA IOTA				202	30.32	48.48		1 00	22.51	AAAA
ATO:	240			202	31.21	6 48.3		าว 1 (10	20.12	AAAA
ATO	M 241				29.12	8 48.9		→	18.04	AAAA
OTA	M 240	и во			28.87		14 61.5	11 1.00	21.09	AAAA
ATO	2.41	09 C	A AR	~ ~ ~ ^	27.56		01 60.7	92 1.00	24.34	AAAA
ATC	OM 24		B AR	704	27.53		59 61.2	06 1.00	27.09 45.73	AAAA
ATC	ъм 24			204	26.25		09 61.1	16 1.0	0 39.82	AAAA
ATO	OM 24		D AR		25.09		:/a bl.c		0 28.40	AAAA
ATO	U1-1		CZ AF	- 701	23.96 23.8		:sn 62.5		0 41.77	AAAA
TA			NH1 AF		22.9	91 50.6	667 61.		0 21.00	AAAA
PΑ	~ ~ /	115 116	NH2 A	3G 304	28.7	94 48.	073 62.4	1 (n 19.45	AAAA
TA				RG 304	29.3	13 48.	087 63.	706 1 (10 19.93	AAAA
	~ ~	418		RG 304	28.1	59 47.	008 PT.	can 1 (າດ 18.70	AAAA AAAA
		419	N A	LA 305	28.0	ino 45.	809 62.		nn 18.26	AAAA
		420	CA A	LA 305	26.9	102 44.		035 1	nn 16.46	AAAA.
	1011	421		LA 305	29.3	311 45.		262 1	nn 19.49	AAAA
	10	422	-	LA 305	29.5	564 44.		005 1	1.92 ר' ממ	AAAA
		423		200	30.	152 44		403]	00 18.99	AAAA
	TOM	2424		206	31.		065 60	.902 1.	00 17.96	AAAA
7		2425		200	32.		. EGA 60	333 1.	00 21.34	AAAA
	TOM .	2426	- -	200	31.		059 59	.014 i	00 16.55	AAAA
	MOT	2427	CD2	4 4 4 4	31.		785 58	.949 1	00 19.37	AAAA
	MOTA	2428		TRP 306	31.		556 57	.878 ¹	.00 17.80	AAAA ,
	MOTA	2429	CE2 CE3	1 1 1 1	32.		610 60	.995 1	.00 19.88 .00 15.95	AAAA
	MOTA	2430	CD1	7 7 4 4	30.		v = 25 60		.00 24.85	/m
	MOTA	2431	NE1	TRP 306	30		1002 57		.00 29.69	• ·- ·- ·
	MOTA	2432	CZ2	TRP 306	; 31		. 701 ht		.00 17.04	
	MOTA	2433	cz3	TRP 306	32		n = 12 5 (00 20.20	,
	MOTA	2434	CH2	TRP 300	5 31		169 6	ل 3.018	.00 21.2	1
	MOTA	2435	_	TRP 30	6 32	159 4	4.726 6		00 18.6	
	MOTE	2436		TRP 30		061 4	6.491 6		າ ດດ 16.8	8
	ATOM	2437		THR 30	. 22	843 4	7.412 0	2 2 2 2	າ ຄຄ 22.0	5
	ATOM	2438		πHR 30	'	579 4	8.885 6	061	1 00 21.5	8
	MOTA	2439	CB	THR 30		218 4	9.132	2.051 4.356	1.00 24.8	6 AAAA
	MOTE	244	nG1	THR 30	· •	3.126	9.857	94	_ •	•
	MOTA	244		THR 30						
	MOTA	544	-	•	•					
				A						

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ATOM	2443	C	THE	307	32.493	47.146	65.187	1.00	17.47	AAAA
ATOM	2444	0	THE	307	33.377	47.142	66.039	1.00	18.94	
ATOM	2445	N	LEU	308	31.216	46.901	65.487	1.00	19.97	AAAA
MOTA	2446	CA	LEU		30.834	46.587	66.866	1.00	22.54	AAAA
MOTA	2447				29.318	46.365	66.989	1.00	21.13	AAAA
MOTA	2448	CG	LEU	308	28.415	47.579	66.751	1.00	22.82	AAAA
MOTA	2449	CD	1 LEU	308	26.937	47.219	67.023		25.01	AAAA
MOTA	2450	CD	2 LEU	308	28.870	48.710	67.685	1.00	29.09	AAAA
MOTA	2451	С	LEU	308	31.578	45.331	67.336		22.98	AAAA
ATOM	2452	. 0	LEU	308	32.056	45.250	68.479		22.27	AAAA
MOTA	2453	. N	ILE	309	31.677	44.342	66.454		22.54	AAAA
MOTA	2454	CA	ILE	309	32.377	43.114	66.801		17.09	AAAA
MOTA	2455	CB	ILE	309	32.318	42.073	65.664		18.12	AAAA ·
ATOM	2456	CG:	2 ILE	309	33.170	40.870	66.033		24.16	AAAA
ATOM	2457	CG:	l ILE	309	30.871	41.655	65.399		18.26	AAAA
ATOM	2458	CD:	l, ILE	309	30.205	40.989			26.57	AAAA
MOTA	2459	С	ILE	309	33.849	43.410	67.067		20.84	AAAA
MOTA	2460	0	ILE	309	34.426	42.905	68.031	1.00	25.20	AAAA
MOTA	2461	N	, TRP	310	34.466	44.223	66.214	1.00	16.86	AAAA
ATOM	2462	.CA	TRP	310	35.888	44.517	66.411	1.00	17.86	AAAA
ATOM	2463	CB	TRP	310	36,439	45.319	65.235	1.00	14.83	AAAA
MOTA	2464	CG	TRP	310	37.879	45.648	65.397	1.00	16.63	AAAA
ATOM	2465		? TRP	310	38.967	44.718	65.560	1.00	18.62	AAAA
ATOM	2466		TRP	310	40.131	45.478	65.799	1.00	25.60	AAAA
ATOM	2467		TRP	310	39.069	43.319	65.529	1.00	24.06	AAAA
ATOM	2468		TRP	310	38.418	46.895	65.533		19.82	AAAA
ATOM	2469		TRP	310	39.768	46.801	65.777		25.84	AAAA
ATOM	2470		TRP	310	41.383	44.887	66.006		26.14	AAAA
MOTA	2471		TRP	310	40.308	42.730	65.735		24.89	AAAA
MOTA	2472 2473		TRP	310	41.452	43.515	65.971		24.96	AAAA
MOTA MOTA	2474	С 0	TRP TRP	310 310	36.112	45.263	67.733		20.86	AAAA
ATOM	2475	N	CYS	311	37.050	44.957	68.476		21.38	AAAA
ATOM	2476	CA	CYS	311	35.242 35.349	46.226 46.971	68.030		24.22	AAAA
ATOM	2477	CB	CYS	311	34.297	48.097	69.280 69.343		27.66	AAAA
ATOM	2478	SG	CYS	311	34.618	49.528	68.253		25.37	AAAA
ATOM	2479	c	CYS	311	35.224	46.042	70.490		27.22 22.95	AAAA AAAA
ATOM	2480	ŏ	CYS	311	35.986	46.180	71.441		25.47	AAAA
ATOM	2481	N	GLU	312	34.284	45.089	70.457		17.03	AAAA
ATOM	2482	CA	GLU	312	34.120	44.129	71.569		22.44	AAAA
ATOM	2483	CB	GLU	312	33.011	43.110	71.280		20.81	AAAA
ATOM	2484	CG	GLU	312	31.856	43.048	72.258		43.65	AAAA
MOTA	2485	CD	GLU	312	32.265	42.971	73.717		29.63	AAAA
MOTA	2486	OE1	GLU	312	33.022	42.059	74.119		38.85	AAAA
ATOM	2487	OE2	GLU	312	31.804	43.844	74.473		53.22	AAAA
ATOM	2488	С	GLI	312	35.395	43.309	71.778		27.47	AAAA
ATOM	2489	0	GL/	312	35.899	43.178	72:895	1.00	22.33	AAAA
ATOM	2490	N	LEL	313	35.899	42.723	70.696	1.00	23.82	AAAA
ATOM	2491	CA	LEU	313	37.101	41.889	70.771	1.00	20.72	AAAA
ATOM	2492	CB	LEU	313	37.380	41.222	69.422	1.00	27.82	AAAA
ATOM	2493	CG	LEU	313	36.403	40.167	68.903	1.00	33.55	AAAA
MOTA	2494		LEU	313	36.839	39.738	67.512	1.00		AAAA
ATOM	2495		LEU	313	36.379	38.981	69.846	1.00	28.20	AAAA
ATOM	2496	C	LEU	313	38.343	42.670	71.181		18.21	AAAA
ATOM	2497	0	LEU	313	39.119	42.205	72.017		21.48	AAAA
ATOM	2498	N	SER	314	38.492	43.848	70.580		19.41	AAAA
ATOM	2499	CA	SER	314	39.627	44.753	70.775		28.26	AAAA
ATOM	2500	CB	SER	314	39.625	45.821	69.663	1.00		AAAA
ATOM	2501	OG	SER	314	40.732	46.696	69.759	1.00		AAAA
MOTA	2502	C	SER	314	39.619	45.429	72.144	1.00		AAAA
ATOM	2503	0	SER	314	40.631	45.969	72.590	1.00		AAAA
MOTA	2504		GLY	315	38.477	45.407	72.806	1.00		AAAA
ATOM	2505	CA	GLY	315	38.393	46.009	74.119	1.00		AAAA
ATOM	2506		GLY	315	38.324	47.518	74.105	1.00		AAAA
ATOM			GLY	315	38.811 37.739	48.178	75.022	1.00		AAAA
MOTA	2508	N .	ARG	316	31.133	48.090	73.065	1.00	21.23	. AAAA

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MOTA	2509	CA	ARG	316	37.631	49.536	73.042	1.00 39.10	AAAA
ATOM	2510	CB	ARG	316	38.347	50.108	71.830	1.00 45.15	AAAA
ATOM	2511	CG	ARG	316	37.722	49.834	70.501	1.00 46.02	AAAA
	2512					50.459		1.00 44.83	
MOTA		CD	ARG	316	38.620		69.449		AAAA
ATOM	2513	NE	ARG	316	39.898	49.767	69.357	1.00 37.91	AAAA
MOTA	2514	CZ	ARG	316	40.945	50.219	68.674	1.00 27.39	AAAA
ATOM	2515	NH1	ARG	316	40.854	51.371	68.034	1.00 50.24	AAAA
MOTA	2516	NH2	ARG	316	42.054	49.493	68.572	1.00 34.51	AAAA
MOTA	2517	С	ARG	316	36.179	49.984	73.058	1.00 35.43	AAAA ·
ATOM	2518	ō	ARG	316	35.292	49.271	72.596	1.00 30.71	AAAA
			GLU	317		51.162			
ATOM	2519	N			35.931		73.612	1.00 34.06	AAAA
ATOM	2520	CA	GLU	317	34.569	51.663	73.671	1.00 37.96	
MOTA	2521	CB	GLU	317	34.481	52.914	74.552	1.00 43.60	AAAA
ATOM	2522	CG	GLU	317	33.961	52.630	75.960	1.00 60.36	AAAA
ATOM	2523	CD	GLU	317	34.768	51.575	76.701	1.00 70.70	AAAA
MOTA	2524	OE1	GLU	317	34.375	51.217	77.832	1.00 76.71	AAAA
ATOM	2525		GLU	317	35.793	51.104	76.162	1.00 78.36	AAAA
ATOM	2526	c	GLU	317	34.068	51.958	72.280	1.00 35.65	
									AAAA
ATOM	2527	0	GLU	317	34.843	52.322	71.390	1.00 32.91	AAAA
ATOM	2528	N	VAL	318	32.767	51.772	72.094	1.00 30.52	AAAA
MOTA	2529	CA	VAL	318	32.138	52.012	70.808	1.00 37.04	AAAA
MOTA	2530	CB	VAL	318	30.877	51.138	70.638	1.00 36.48	AAAA
ATOM	2531	CG1	VAL	318	30.278	51.366	59.268	1.00 40.43	AAAA
ATOM	2532	CG2	VAL	318	31.222	49.674	70.846	1.00 33.75	AAAA
ATOM	2533	C	VAL	318	31.719	53.465	70.737	1.00 28.96	AAAA
ATOM	2534	ŏ	VAL	318	30.930	53.915	71.556	1.00 33.56	AAAA
	2535				32.258	54.229	69.773		
MOTA		N	PRO	319				1.00 29.20	AAAA
MOTA	2536	CD	PRO	319	33.243	53.924	68.726	1.00 31.62	AAAA
MOTA	2537	CA	PRO	319	31.858	55.637	69.684	1.00 28.99	AAAA
MOTA	2538	CB	PRO	319	32.709	56.154	68.528	1.00 32.17	AAAA
ATOM	2539	CG	PRO	319	32.850	54.926	67.664	1.00 41.36	AAAA
ATOM	2540	С	PRO	319	30.365	55.680	69.377	1.00 36.95	AAAA
ATOM	2541	0	PRO	319	29.847	54.795	68.695	1.00 32.86	AAAA
ATOM	2542	N	GLU	320	29.646	56.683	69.855	1.00 34.61	AAAA
ATOM	2543	CA	GLU	320	28.230	56.657	69.544	1.00 35.13	AAAA
	2544		GLU	320	27.419	57.416	70.595	1.00 52.97	AAAA
ATOM		CB					70.738		
MOTA	2545	CG	GLU	320	27.751	58.875		1.00 56.06	AAAA
MOTA	2546	CD	GLU	320	26.822	59.558	71.721	1.00 65.58	AAAA
MOTA	2547	OE1		320	25.604	59.619	71.444	1.00 64.27	AAAA
MOTA	2548	OE2	GLU	320	27.306	60.022	72.775	1.00 72.99	AAAA
ATOM	2549	C	GLU	320	27.943	57.192	68.153	1.00 35.13	AAAA
MOTA	2550	0	GLU	320	26.916	56.879	67.565	1.00 37.43	AAAA
ATOM	2551	N	LYS	321	28.880	57.953	67.604	1.00 28.22	AAAA
ATOM	2552	CA	LYS	321	28.700	58.555	66.289	1.00 36.58	AAAA
	2553	CB	LYS	321	28.666	60.071	66.454	1.00 44.87	AAAA
ATOM									
ATOM	2554	CG	LYS	321	29.987	60.606	67.023	1.00 55.73	'AAA
MOTA	2555	CD	LYS	321	30.305	60.020	68.410	1.00 57.27	:AAA:
ATOM	2556	CE	LYS	321	31.733	60.310	68.840	1.00 54.59	AAA.
ATOM	2557	NZ	LYS	321	32.024	61.774	68.848	1.00 67.47	AAAA
ATOM	2558	С	LYS	321	29.823	58.211	65.315	1.00 34.44	AAAA
ATOM	2559	0	LYS	321	30.912	57.818	65.731	1.00 33.83	AAAA
MOTA	2560		LEU	322	29.549	58.354	64.019	1.00 30.21	AAAA
MOTA	2561		LEU	322	30.575	58.135	62.998	1.00 29.45	AAAA
					29.966	57.677	61.677	1.00 32.21	AAAA
ATOM	2562		LEU	322					
ATOM	2563		LEU	322	29.240	56.338	61.651	1.00 38.94	AAAA
ATOM	2564	CD1		322	29.008	55.977	60.186	1.00 38.44	AAAA
ATOM	2565	CD2	LEU	322	30.072	55.261	62.337	1.00 42.11	AAAA
MOTA	2566	С	LEU	322	31.228	59.503	62.783	1.00 33.28	AAAA
ATOM	2567		LEU	322	30.544	60.519	62.872	1.00 31.45	AAAA
ATOM	2568		ASN	323	32.533	59.539	62.519	1.00 34.38	AAAA
ATOM	2569		ASN	323	33.208	60.824	62.294	1.00 36.53	AAAA
			ASN	323	34.701	60.737	62.600	1.00 42.85	AAAA
MOTA	2570								_
ATOM	2571		ASN	323	35.484	60.081	61.480	1.00 50.51	AAAA
ATOM ·		OD1		323	35.215	58.942	61.109	1.00 51.23	AAAA
ATOM .	2573	ND2		323	36.455	60.807	60.928	1.00 60.23	AAAA
MOTA	2574	.C	ASN	323	33.027	61.171	60.822	1.00 34.69	AAAA
		•							•

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ATOM	2579	5 0	ASN	323	32.429	60.395	60.075	1.00 34.06	AAAA
MOTA	2576	5 N	ASN	324	33.551	62.317	60.390		AAAA
ATOM	2577				33.385				
	2578								AAAA
MOTA					33.868				AAAA
ATOM	2579				32.974		59.450	1.00 45.77	AAAA
ATOM	2580	OD (1 ASN	324	31.765	65.189	59.206	1.00 39.55	AAAA
MOTA	2581	ND	2 ASN	324	33.555				AAAA
ATOM	2582		ASN		34.047				
								1.00 24.87	AAAA
MOTA	2583		ASN		33.451			1.00 31.91	AAAA
ATOM	2584	N	LYS	325	35.276	61.405	58.250	1.00 27.73	AAAA
MOTA	2585	CA	LYS	325	- 35.991	60.538	57.333	1.00 29.55	AAAA
MOTA	2586	CB	LYS	325	37.351	60.182		1.00 37.43	AAAA
ATOM	2587			325	38.250				-
					39.684			1.00 44.84	AAAA
MOTA	2588			325				1.00 50.89	AAAA
ATOM	2589		LYS	325	40.191		57.561	1.00 54.82	AAAA
ATOM	2590	NZ	LYS	325	41.621	60.980	57.969	1.00 65.70	AAAA
ATOM	2591	С	LYS	325	35.161	59.279	57.078	1.00 27.99	AAAA
ATOM	2592		LYS	325	35.016			1.00 31.80	
									AAAA
ATOM	2593		ALA	326	34.602	58.721	58.142	1.00 26.07	AAAA
ATOM	2594		ALA	326	33.781	57.506	58.030	1.00 24.38	AAAA
MOTA	2595	CB	ALA	326	33.470	56.982	59.428	1.00 27.34	AAAA
MOTA	2596	С	ALA	326	32.478	57.709	57.231	1.00 25.78	AAAA
ATOM	2597	0	ALA	326	32.131	56.890	56.369	1.00 27.37	AAAA
ATOM	2598	N	LYS	327	31.749	58.790	57.496	1.00 27.31	
		CA							AAAA
ATOM	2599		LYS	327	30.502	59.027	56.758	1.00 28.56	AAAA
ATOM	2600	CB	LYS	327	29.759	60.251	57.313	1.00 28.87	AAAA
ATOM	2601	CG	LYS	32 7	29.491	60.209	58.812	1.00 36.72	AAAA
ATOM	2602	CD	LYS	327	· 28.643	61.407	59.255	1.00 40.34	AAAA
ATOM	2603	CE	LYS	327	28.645	61.594	60.769	1.00 38.91	AAAA
ATOM-	2604	NZ	LYS	327	28.163	60.429	61.556	1.00 47.67	
ATOM	2605	C	LYS		30.792			-	AAAA
						59.244	55.269	1.00 29.15	AAAA
MOTA	2606	0	LYS	327	30.097	58.719	54.393	1.00 27.76	AAAA
ATOM	2607	N	GLU	328	31.829	60.015	54.972	1.00 31.59	AAAA
ATOM	2608	CA	GLU	328	32.167	60.265	53.581	1.00 28.93	AAAA
MOTA	2609	CB	GLU	328	33.257	61.332	53.515	1.00 32.30	AAAA
ATOM	2610	CG	GLU	328	32.745	62.652	54.067	1.00 47.50	
ATOM	2611	CD	GLU	328					AAAA
					33.764	63.772	54.032	1.00 46.67	AAAA
ATOM	2612	OE1		328	34.325	64.037	52.951	1.00 56.88	AAAA
MOTA	2613	OE2	GLU	328	33.984	64.402	55.087	1.00 42.24	AAAA
ATOM	.2614	С	GLU	328	32.575	58.975	52.871	1.00 30.46	AAAA
ATOM	2615	0	GLU	328	32.226	58.753	51.704	1.00 26.29	AAAA
ATOM	2616	N	LEU	329	33.292	58.112	53.584	1.00 24.93	AAAA
MOTA	2617	CA	LEU	329	33.701	56.828	53.017	1.00 24.80	
	2618	CB							AAAA
ATOM			LEU	329	34.478	56.003	54.053	1.00 25.70	AAAA
ATOM	2619	CG	LEU	329	34.730	54.522	53.703	1.00 19.71	AAAA
ATOM	2620	CD1	LEU	329	5.569	54.413	52.430	1.00 25.26	AAAA
MOTA	2621	CD2	LEU	329	55.412	53.833	54.863	1.00 24.73	AAAA
MOTA	2622	С	LEU	329	.2.443	56.059	52.603	1.00 23.50	AAAA
ATOM	2623	0	LEU	329	32.310	55.650	51.453	1.00 25.60	
ATOM		N							AAAA
	2624		LEU	330	31.516	55.881	53.539	1.00 23.02	AAAA
ATOM	2625	CA	LEU	330	30.289	55.145	53.242	1.00 23.85	AAAA
MOTA	2626	CB	LEU	330	29.414	55.030	54.484	1.00 21.74	AAAA
ATOM	2627	CG	LEU	330	30.039	54.252	55.642	1.00 25.29	AAAA
MOTA	2628	CD1	LEU	330	28.984	54.053	56.724	1.00 30.58	AAAA
ATOM	2629		LEU	330	30.538	52.905			
							55.168	1.00 22.44	AAAA
ATOM	2630	S.	LEU	330	29.491	55.769	52.113	1.00 26.94	AAAA
MOTA	2631	Ö	LEU	330	28.968	55.060	51.252	1.00 26.65	AAAA
ATOM	2632	N	LYS	331	29.404	57.097	52.111	1.00 30.82	AAAA
ATOM	2633	CA	LYS	331	28.667	57.795	51.066	1.00 29.53	AAAA
ATOM	2634	CB	LYS	331	28.537	59.292	51.407	1.00 29.67	AAAA
ATOM	2635	CG	LYS	331	27.814	59.567	52.714	1.00 36.06	AAAA
ATOM	2636	CD	LYS	331	27.688	61.055	52.990	1.00 42.75	AAAA
ATOM	2637	CE	LYS	331	26.828	61.737	51.939	1.00 53.98	AAAA
MOTA	2638	NZ	LYS	331	26.634	63.185	52.234	1.00 67.57	AAAA
ATOM	2639	C	LYS	331	29.315	57.628	49.692	1.00 30.07	AAAA
	2640	0	•		28.634				AAAA
ATOM	2040	9	LYS	331	20.034	57.759	48.672	1.00 36.20	WWW.

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MOTA	2641	N	SER	332	30.608	57.305	49.657	1.00 30.08	AAAA
ATOM	2642	CA	SER	332	31.322	57.153	48.385	1.00 33.35	AAAA
				.332			48.590		
ATOM	2643	CB	SER		32.834	57.312		1.00 40.36	AAAA
MOTA	2644	OG	SER	332	33.396	56.169	49.219	1.00 34.04	AAAA
ATOM	2645	С	SER	332	31.061	55.821	47.693	1.00 37.72	AAAA
ATOM	2646	0	SER	332	31.354	55.661	46.507	1.00 30.78	AAAA
ATOM	2647	N	ILE	333	30.521	54.865	48.440	1.00 30.61	AAAA
								1.00 37.59	
MOTA	2648	CA	ILE	333	30.219	53.547	47.899	· · · · · · · · · · · · · · · · · · ·	` AAAA
MOTA.	2649	CB	ILE	- 333	29.901	52.551	49.022	1.00 33.59	AAAA
ATOM	2650	CG2	. ILE	333	29.738	51.146	48.442	1.00 37.05	AAAA
MOTA	2651	CG1	ILE	333	31.015	52.564	50.065	1.00 38.95	AAAA
MOTA	2652	CD1	ILE	. 333	30.706	51.727	51.282	1.00 46.83	AAAA
ATOM	2653	C	ILE	333 .	28.990	53.620	46.998	1.00 43.41	AAAA
				333	27.889	53.876	47.479		
ATOM	2654	0	ILE					1.00 46.24	AAAA
ATOM	2655	N	ASP	334	29.158	53.423	45.696	1.00 47.97	AAAA
ATOM	2656	ÇA	ASP	334	27.976	53.447	44.847	1.00 53.47	- AAAA
ATOM	2657	CB	ASP	334	28.333	53.535	43.358	1.00 61.52	AAAA
ATOM	2658	CG	ASP	334	29.223	52.406	42.897	1.00 64.75	AAAA
ATOM	2659		ASP	334	29.379	52.248	41.666	1.00 66.93	AAAA
ATOM	2660		ASP	334	29.779	51.691	43.758	1.00 65.93	
ATOM	2661	C	ASP	334	27.248	52.144	45.161	1.00 51.83	AAAA
ATOM	2662	0	ASP	334	27.626	51.067	44.699	1.00 46.80	AAAA
MOTA	2663	N	PHE	335	26.215	52.249	45.986	1.00 54.96	aaaa
ATOM	2664	CA	PHE	335	25.455	51.080	46.392	1.00 50.60	AAAA
ATOM	2665	CB	PHE	335	25.413	51.003	47.920	1.00 39.55	AAAA
ATOM	2666	CG	PHE	335	24.380	50.054	48.440	1.00 37.98	AAAA
ATOM	2667		PHE	335	24.389	48.715	48.054	1.00 46.72	AAAA
	2668		PHE	335	23.362	50.506	49.262	1.00 34.23	
ATOM									AAAA
MOTA	2669		PHE	335	23.389	47.842	48.478	1.00 49.80	AAAA
ATOM	2670	CE2	PHE	335	22.361	49.644	49.689	1.00 48.51	AAAA
MOTA	2671	CZ	PHE	335	22.373	48.309	49.296	1.00 40.44	AAAA
ATOM	2672	С	PHE	335	24.033	51.000	45.839	1.00 54.52	AAAA
ATOM	2673	0	PHE	335	23.603	49.939	45.379	1.00 59.24	AAAA
ATOM	2674	N	GLU	336	23.302	52.108	45.888	1.00 50.94	AAAA
ATOM	2675	CA	GLU	336	21.923	52.119	45.406	1.00 57.05	AAAA
MOTA	2676	CB	GLU	336	21.853	51.751	43.924	1.00 60.27	AAAA
MOTA	2677	CG	GLU	336	20.430	51.627	43.422	1.00 68.55	AAAA
MOTA	2678	CD	GLU	336	20.352	51.126	42.001	1.00 80.03	AAAA
ATOM	2679	OE1	GLU	336	20.860	50.013	41.735	1.00 84.64	AAAA
ATOM	2680	OE2	GLU	336	19.777	51.841	41.153	1.00 80.68	AAAA
MOTA	2681	С	GLU	336	21.065	51.135	46.201	1.00 55.73	AAAA
ATOM	2682	ō	GLU	336	21.219	49.917	46.089	1.00 51.33	AAAA
ATOM	2683	N	GLU	337	20.151	51.679	46.992	1.00 49.54	AAAA
				337	19.267		47.821	1.00 48.19	AAAA
ATOM	2684	CA	GLU			50.880			
MOTA	2685	CB	GLU	337	18.510	51.822	48.764	1.00 47.73	AAAA
MOTA	2686	CG	GLU	337	18.084	51.205	50.077	1.00 55.69	AAAA
ATOM	2687	CD	GLU	. 337	19.269	50.720	50.904	1.00 50.17	AAAA
ATOM	2688	OE1	GLU	337	20.111	51.548	51.345	1.00 36.03	AAAA
ATOM	2689	OE2	GLU	337	19.358	49.494	51.105	1.00 51.25	AAAA
ATOM	2690	C	GLU	337	18.294	50.083	46.936	1.00 49.13	AAAA
	2691	Ö	GLU	337	17.816	50.588	45.916	1.00 48.61	AAAA
MOTA									
ATOM	2692	N	PHE	338	18.015	48.837	47.313	1.00 48.15	AAAA
ATOM	2693	CA	PHE	338	17.092	48.000	46.547	1.00 48.12	AAAA
MOTA	2694	CB	PHE	338	16.870	46.658	47.249	1.00.54.54	AAAA
ATOM	2695	CG	PHE	338	15.883	45.777	46.548	1.00 57.22	AAAA
ATOM:	2696	CD1		338	16.115	45.366	45.243	1.00 60.01	AAAA
	2697	CD2		338	14.699	45.398	47.171	1.00 55.04	AAAA
MOTA					15.185		44.566	1.00 60.84	AAAA
MOTA	2698	CE1		338		44.597			
ATOM	2699	CE2		338	13.758	44.624	46.497	1.00 59.41	AAAA
MOTA	2700	CZ	PHE	338	14.002	44.224	45.189	1.00 57.18	AAAA
MOTA	2701	С	PHE	338	15.755	48.714	46.380	1.00 45.46	AAAA
ATOM	2702	0	PHE	338	15.274	48.900	45.263	1.00 51.11.	AAAA
ATOM	2703	N	ASP	339	15.154	49.098	47.501	1.00 40.38	AAAA .
			ASP	339	13.890	49.820	47.488	1.00 49.97	AAAA
ATOM.	2704			339	13.270		48.886	1.00 53.23	AAAA
MOTA	2705	CB	ASP			49.821		1.00 57.40	
ATOM	2706	CG	ASP	339	12.000	50.659	48.968	1.UU 3/.4U	AAAA .

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Figure 16-42

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ATOM	2707	OD	1 ASP	339	12.039	51.858	48.616	1.00 53.79	AAAA
ATOM	2708		2 ASP		10.963	50.118		1.00 51.15	
									AAAA
ATOM	2709	С	ASP		14.215			1.00 55.06	AAAA
MOTA	2710	0	ASP		14.994	51.922		1.00 56.47	AAAA
MOTA	2711	N	ASP	340	13.623	51.708	45.978	1.00 58.46	AAAA
ATOM	2712	CA	ASP	340	13.874	53.059	45.484	1.00 67.72	AAAA
ATOM	2713	CB	ASP		12.683	53.559	44.664	1.00 71.52	
	2714	CG			12.611	_			AAAA
MOTA						52.913	43.295	1.00 79.72	AAAA
MOTA	2715		1 ASP		12.528	51.667		1.00 86.74	AAAA
ATOM	2716	OD:	2 ASP	340	12.640	53.655	42.288	1.00 83.40	AAAA
MOTA	2717	C	ASP	340	14.209	54.072	46.572	1.00 69.65	AAAA
MOTA	2718	0	ASP	340	15.204	54.794	46.463	1.00 70.13	AAAA
ATOM	2719	Ň	GLU		13.392	54.130	47.620		
								1.00 67.11	AAAA
MOTA	2720	CA	GLU		13.668	55.077	48.689	1.00 67.87	AAAA
ATOM	2721	CB	GLU		13.195	56.478	48.278	1.00 74.87	AAAA
MOTA	2722	CG	GLU	341	13.502	57.576	49.298	1.00 82.72	AAAA
ATOM	2723	CD	GLU	341	13.162	58.974	48.790	1.00 90.80	AAAA
ATOM	2724		1 GLU		11.988	59.215	48.431	1.00 90.38	
									AAAA
MOTA	2725		2 GLU		14.072	59.835	48.752	1.00 93.36	AAAA
MOTA	2726	C	GLU	341	13.101	54.719	50.058	1.00 60.22	AAAA
ATOM	2727	0	GLU	341	11.929	54.955	50.347	1.00 58.81	ААЛА
ATOM	2728	N	VAL	342	13.956	54.144	50.897	1.00 57.28	AAAA
ATOM	2729	CA	VAL	342	13.594	53.781	52.262	1.00 52.09	AAAA
ATOM	2730	CB	VAL	342	14.195	52.419	52.669	1.00 53.17	
	2731		VAL		13.730				AAAA
MOTA				342		52.042	54.070	1.00 46.16	AAAA
ATOM	2732		VAL	342	13.815	51.356	51.663	1.00 59.09	AAAA
ATOM	2733	C	VAL	342	14.263	54.843	53.124	1.00 53.31	AAAA
ATOM	2734	0	VAL	342	13.763	55.230	54.185	1.00 57.79	AAAA
ATOM	2735	N	ASP	343	15.398	55.306	52.610	1.00 46.24	AAAA
ATOM	2736	CA	ASP	343	16.268	56.289	53.243	1.00 42.60	AAAA
ATOM	2737	CB	ASP	343	15.521	57.510	53.781	1.00 43.88	
									AAAA
ATOM	2738	CG	ASP	343	16.480	58.581	54.290	1.00 46.82	AAAA
ATOM	2739		. ASP	343	16.028	59.581	54.887	1.00 46.16	AAAA
MOTA	2740	OD2	ASP	343 .	17.700	58.414	54.075	1.00 33.01	AAAA
ATOM	2741	С	ASP	343	17.012	55.636	54.395	1.00 35.45	AAAA
ATOM	2742	0	ASP	343	16.487	55.480	55.502	1.00 29.39	AAAA
ATOM	2743	N	ARG	344	18.247	55.249	54.124	1.00 30.51	AAAA
ATOM	2744	CA	ARG	344	19.059				
						54.613	55.140	1.00 29.43	AAAA
MOTA	2745	CB	ARG	344	19.736	53.377	54.561	1.00 30.10	AAAA
ATOM	2746	CG	ARG	344	18.803	52.258	54.180	1.00 33.95	AAAA
ATOM	2747	CD	ARG	344	17.981	51.770	55.365	1.00 20.92	AAAA
ATOM	2748	NE	ARG	344	17.120	50.673	54.936	1.00 29.72	AAAA
MOTA	2749	CZ	ARG	344	16.110	50.176	55.639	1.00 29.13	AAAA
ATOM	2750		ARG	344	15.805	50.668	56.835	1.00 29.63	AAAA
	2751								
MOTA			ARG	344	15.379	49.198	55.120	1.00 27.19	AAAA:
ATOM	2752	С	ARG	344	20.116	55.769	55.660	1.00 34.31	AAAA
ATOM	2753	0	ARG	344	21.005	5557	56.391	1.00 29.09	AAAA
ATOM	2754	N	SER	345	20.011	56. 45	55.294	1.00 28.34	AAAA
ATOM	2755	CA	SER	345	20.999	57.839	55.715	1.00 30.95	AAAA
ATOM	2756	CB	SER	345	20.669	59.199	55.109	1.00 29.56	AAAA
	2757								
ATOM		OG	SER	345	19.429	59.648	55.610	1.00 29.38	AAAA
MOTA	2758	С	SER	345	21.137	57.988	57.230	1.00 30.92	AAAA
MOTA	2759	0	SER	345	22.155	58.488	57.718	1.00 31.15	AAAA
MOTA	2760	N	TYR	346	20.116	57.576	57.975	1.00 25.64	AAAA
ATOM	2761	CA	TYR	346	20.158	57.659	59.433	1.00 26.81	AAAA
MOTA	2762	СВ	TYR	346	18.823	57.189	60.006	1.00 34.41	AAAA
ATOM	2763	CG	TYR	346	18.529	55.723	59.716	1.00 27.35	AAAA
MOTA	2764		TYR	346	19.003	54.708	60.556	1.00 24.87	AAAA
ATOM	2765	CE1	TYR	346	18.744	53.352	60.278	1.00 28.05	AAAA
MOTA	2766	CD2	TYR	346	17.795	55.358	58.588	1.00 27.70	AAAA
ATOM	2767		TYR	346	17.533	54.008	58.297	1.00 26.59	AAAA
					18.008				
ATOM	2768	CZ	TYR	346		53.015	59.145	1.00 33.75	AAAA
ATOM	2769	OH	TYR	346	17.737	51.691	58.855	1.00 26.06	AAAA
ATOM .	2770	С	TYR	346	21.277	56.766	59.977	1.00 25.57	, AAAA
ATOM	2771	0	TYR	346	21.769	56.970	61.085	1.00 28.07	AAAA
ATOM	2772	N .	MET	347	21.666	55.761	59.198	1.00 29.08	AAAA
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MOTA	2773	CA	MET	347	22.720	54.837	59.622	1.00 24.19	AAAA
ATOM	2774	CB	MET	347	22.844	53.678	58.628	1.00 24.87	AAAA
					21.609	52.806	58.543	1.00 23.66	AAAA
MOTA	2775	CG	MET	347					
ATOM	2776	SD	MET	347	21.780	51.503	57.267	1.00 27.02	AAAA
ATOM	2777	CE	MET	347	22.115	52.375	55.896	1.00 37.69	AAAA
		c	MET	347	24.054	55.540	59.737	1.00 29.45	AAAA
MOTA	2778								
ATOM	2779	0	MET	347	24.937	55.092	60.479	1.00 28.08	AAAA
ATOM	2780	N	LEU	348	24.188	56.650	59.007	1.00 23.71	AAAA
						57.446	58.998	1.00 34.11	AAAA
MOTA	2781	CA	LEU	348	25.418				
ATOM	2782	CB	LEU	348	25.463	58.351	57.757	1.00 25.37	AAAA
ATOM	2783	CG	LEU	348	25.320	57.785	56.344	1.00 30.38	AAAA
					25.307	58.944	55.340	1.00 27.44	AAAA
ATOM	2784		LEU	348					
ATOM	2785	CD2	LEU	348	26.459	56.814	56.041	1.00 36.44	AAAA
	2786	С	LEU	348	25.507	58.332	60.237	1.00 36.09	AAAA
ATOM						58.894	60.539	1.00 33.30	AAAA
ATOM	2787	0	LEU	348	26.561				
ATOM	2788	N	GLU	349	24.394	58.445	60.953	1.00 30.51	AAAA
	2789	CA	GLU	349	24.313	59.292	62.136	1.00 35.53	AAAA
MOTA						59.896	62.217	1.00 31.35	AAAA
ATOM	2790	CB	GLU	349	22.908				
ATOM	2791	CG	GLU	349	 22.518	60.717	61.006	1.00 29.09	AAAA
ATOM	2792	CD	GLU	349	23.481	61.859	60.746	1.00 31.78	AAAA
					23.937	62.476	61.730	1.00 30.98	AAAA
MOTA	2793		GLU	349					
MOTA	2794	OE2	GLU	349	23.766	62.155	59.569	1.00 30.67	AAAA
MOTA	2795	С	GLU	349	24.663	58.633	63.471	1.00 38.48	AAAA
				349	24.727	59.303	64.502	1.00 40.12	AAAA
MOTA	2796	0	GLU						AAAA
MOTA	2797	N	THR	350	24.878	57.326	63.461	1.00 33.58	
ATOM	2798	CA	THR	350	25.221	56.612	64.681	1.00 29.74	AAAA
		CB	THR	350	23.992	56.363	65.559	1.00 35.91	AAAA
MOTA	2799							1.00 45.03	AAAA
ATOM	2800	OG1	THR	350	23.421	57.615	65.952		
ATOM	2801	CG2	THR	350	24.382	55.586	66.806	1.00 49.48	AAAA
	2802	С	THR	350	25.821	55.267	64.330	1.00 30.63	AAAA
ATOM						54.709	63.274	1.00 26.62	AAAA
ATOM	2803	0	THR	350	25.535				
ATOM	2804	N	LEU	351	26.644	54.740	65.225	1.00 29.07	AAAA
ATOM	2805	CA	LEU	351	27.271	53.461	64.972	1.00 24.59	AAAA
				351	28.584	53.367	65.757	1.00 29.91	AAAA
MOTA	2806	CB	LEU					1.00 39.62	AAAA
MOTA	2807	CG	LEU	351	29.591	52.327	65.267		
ATOM	2808	CD1	LEU	351	30.887	52.467	66.039	1.00 37.09	AAAA
			LEU	351	29.024	50.935	65.415	1.00 54.03	AAAA
MOTA	2809						65.377	1.00 29.71	AAAA
MOTA	2810	C	LEU	351	26.314	52.336			
MOTA	2811	0	LEU	351	26.130	51.364	64.641	1.00 30.53	AAAA
	2812	N	LYS	352	25.697	52.481	66.543	1.00 28.64	AAAA
ATOM					24.763	51.479	67.061	1.00 32.72	AAAA
ATOM	2813	CA	LYS	352					AAAA
ATOM	2814	CB	LYS	352	24.913	51.381	68.581	1.00 27.37	
ATOM	2815	CG	LYS	352	26.230	50.787	69.034	1.00 43.48	AAAA
			LYS	352	26.536	51.068	70.504	1.00 46.77	AAAA
MOTA	2816	CD					71.451	1.00 51.52	AAAA
ATOM	2817	CE	LYS	352	25.484	50.538			
A OM	2818	NZ	LYS	352	25.850	50.859	72.866	1.00 62.08	AAAA
	2819	C	LYS	352	23.330	51.856	66.731	1.00 32.49	AAAA
A . OM					22.953	53.010	66.882	1.00 31.90	AAAA
A 1 OM	2820	0	LYS	352				1.00 31.44	AAAA
ATOM	2821	N	ASP	353	22.525	50.916	66.244		
ATOM	2822	CA	ASP	353	21.136	51.286	66.012	1.00 26.50	AAAA
				353	20.543	50.635	64.746	1.00 50.09	AAAA
MOTA	2823	CB	ASP				64.604	.1.00 52.79	AAAA
MOTA	2824	CG	ASP	353	20.880	49.176		. 1.00 32.79	
ATOM	2825	OD1	ASP	353	21.980	48.861	64.109	1.00 58.55	AAAA
				353	20.040	48.339	64.984	1.00 73.19	AAAA
ATOM	2826		ASP					1.00 26.41	AAAA
MOTA	2827	С	ASP	353	20.328	50.930	67.257	1,00 20.41	
ATOM	2828	0	ASP	353	20.806	50.214	68.136	1.00 25.73	AAAA
			PRO	354	19.118	51.481	67.385	1.00 30.12	AAAA
MOTA	2829	N					66.495	1.00 35.38	AAAA
ATOM	2830	CD	PRO	354	18.428	52.429			
ATOM	2831	CA	PRO	354	18.276	51.190	68.547	1.00 34.02	AAAA
		CB	PRO	354	17.091	52.129	68.340	1.00 32.25	AAAA
ATOM	2832						66.833	1.00 44.48	AAAA
ATCM	2833	CG	PRO	354	16.974	52.139			
ATOM	2834	С	PRO	354	17.838	49.736	68.512	1.00 34.00	AAAA
		Õ	PRO	354	17.829	49.111	67.452	1.00 28.28	AAAA
ATOM	2835				17.484	49.190	69.664	1.00 23.89	AAAA
ATOM ·	2836	N	TRP	355				1.00 33.84	AAAA
ATOM	2837	CA	TRP	355.	17.010	47.818	69.669		
	2838	CB	TRP	355	16.653	47.363	71.076	1.00 33.84	AAAA
ATOM	2000			٠.		•	-		•
		-							

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Figure 16-44

ATOM 2840 CDZ TRP 355										
ATOM 2840 CDZ TRP 355	MOTA	2839	CG	TRP	355	17.844	46.946	. 71.832	1.00 49.97	AAAA
AROM 2841 CEZ TRP 355	ATOM	2840	CD2	TRP	355	18.364	45.622	71.905	1 00 46 60	
ATOM 2842 CE3 TRP 2555 17.931 44.386 71.419 1.00 46.90 AAAA ATOM 2843 CD1 TRP 355 18.723 47.746 72.507 1.00 56.10 56.10 AAAA ATOM 2844 NE1 TRP 355 18.723 47.746 72.897 1.00 56.07 AAAA ATOM 2846 C23 TRP 355 19.765 46.991 72.897 1.00 55.25 AAAA ATOM 2846 C23 TRP 355 19.867 44.552 72.897 1.00 55.25 AAAA ATOM 2846 C2 TRP 355 19.867 43.356 72.405 1.00 50.68 AAAA ATOM 2848 C TRP 355 19.867 43.356 72.405 1.00 50.68 AAAA ATOM 2848 C TRP 355 15.789 47.712 68.776 1.00 30.312 AAAA ATOM 2851 CA ARG 356 15.547 46.508 68.263 1.00 29.41 AAAA ATOM 2851 CA ARG 356 14.413 46.237 67.387 1.00 23.96 AAAA ATOM 2851 CA ARG 356 14.413 46.237 67.387 1.00 23.96 AAAA ATOM 2852 CB ARG 356 15.507 47.385 65.393 1.00 23.96 AAAA ATOM 2853 CC ARG 356 16.291 47.212 64.108 1.00 29.91 AAAA ATOM 2855 C ARG 356 16.291 47.212 64.108 1.00 29.92 ATOM 2856 C2 ARG 356 16.833 48.503 63.666 1.00 24.73 AAAA ATOM 2857 NH1 ARG 356 AAAA ATOM 2858 NH2 ARG 356 18.209 47.616 62.066 1.00 22.15 AAAA ATOM 2858 NH2 ARG 356 18.209 47.616 62.066 1.00 22.15 AAAA ATOM 2858 CA GLY 357 11.338 49.891 62.418 1.00 22.99 AAAA ATOM 2868 O ARG 356 13.785 43.925 67.189 1.00 22.55 AAAA ATOM 2868 O GLY 357 11.338 49.891 62.418 1.00 22.55 AAAA ATOM 2868 O GLY 357 11.338 49.891 62.418 1.00 22.55 AAAA ATOM 2868 O GLY 357 11.338 49.891 62.418 1.00 22.57 AAAA ATOM 2868 O GLY 357 11.338 49.891 62.418 1.00 22.57 AAAA ATOM 2868 O GLY 357 10.536 44.330 68.619 1.00 22.15 AAAA ATOM 2868 O GLY 357 10.536 44.330 68.619 1.00 22.15 AAAA ATOM 2868 O GLY 357 10.536 44.330 68.619 1.00 22.15 AAAA ATOM 2868 O GLY 358 9.188 8.00 41.274 70.20 1.00 23.91 AAAA ATOM 2868 O GLY 358 9.188 8.00 41.274 70.20 1.00 22.99 AAAA ATOM 2868 O GLY 358 9.626 40.757 71.663 1.00 22.97 AAAA ATOM 2868 O GLY 358 9.397 7.71.163 1.00 29.97 AAAA ATOM 2868 O GLY 358 9.399 7.701 1.366 1.00 22.97 AAAA ATOM 2868 O GLY 358 9.466 6.672 70.10 1.00 29.97 AAAA ATOM 2870 C G GLU 359 7.804 39.327 72.299 1.00 45.44 4.304 ATOM 2870 C G GLU 359 7.804 39.327 72.299 1.00 4.304 ATOM 2870 C G GLU 359 7.804 39.327 72.299 1.00 4.304 ATOM 2870 C										
ATOM 2843 CD1 TRP 355 18.723 47.746 72.507 1.00 56.07 AAAA ATOM 2845 CZ2 TRP 355 19.765 46.991 72.997 1.00 55.05 AAAA ATOM 2846 CZ3 TRP 355 18.696 43.267 71.674 1.00 50.74 AAAA ATOM 2847 CH2 TRP 355 19.867 43.267 71.674 1.00 50.74 AAAA ATOM 2848 C TRP 355 15.789 47.712 68.776 1.00 50.74 AAAA ATOM 2848 C TRP 355 15.789 47.712 68.776 1.00 23.92 41 AAAA ATOM 2849 O TRP 355 15.789 47.712 68.250 1.00 23.90 AAAA ATOM 2851 CA ARG 356 15.547 46.508 68.263 1.00 23.90 AAAA ATOM 2851 CA ARG 356 14.892 46.096 65.995 1.00 23.96 AAAA ATOM 2852 CB ARG 356 14.892 46.096 65.995 1.00 22.66 AAAA ATOM 2854 CD ARG 356 16.291 47.212 64.096 65.995 1.00 22.66 AAAA ATOM 2855 NE ARG 356 16.291 47.212 64.108 1.00 22.96 AAAA ATOM 2855 NE ARG 356 16.291 47.212 64.108 1.00 22.96 AAAA ATOM 2858 NH2 ARG 356 16.291 47.212 64.108 1.00 22.96 AAAA ATOM 2855 NE ARG 356 16.291 47.212 64.108 1.00 22.96 AAAA ATOM 2856 CZ ARG 356 17.733 48.668 62.724 1.00 23.57 AAAA ATOM 2857 NH1 ARG 356 18.209 47.616 62.066 1.00 22.15 AAAA ATOM 2858 NH2 ARG 356 18.153 49.891 62.724 1.00 22.55 AAAA ATOM 2859 C ARG 356 13.785 49.891 62.048 10.02 22.55 AAAA ATOM 2859 C ARG 356 13.785 49.891 62.048 10.02 22.55 AAAA ATOM 2860 O ARG 356 13.785 43.925 67.189 1.00 22.55 AAAA ATOM 2861 N GLY 357 12.631 44.993 69.085 1.00 22.55 AAAA ATOM 2865 N GLY 357 12.631 44.993 69.085 1.00 22.25 AAAA ATOM 2866 C A GLY 357 12.631 44.993 69.085 1.00 22.25 AAAA ATOM 2866 C A GLY 357 12.631 44.993 69.085 1.00 22.25 AAAA ATOM 2867 C GLY 358 88.800 41.274 70.920 1.00 22.25 AAAA ATOM 2866 C A GLY 357 10.536 44.330 68.69 1.00 22.25 AAAA ATOM 2867 C GLY 358 8.800 41.274 70.920 1.00 20.93 AAAA ATOM 2867 C GLY 358 8.800 41.274 70.920 1.00 20.93 AAAA ATOM 2868 C GLY 358 8.800 41.274 70.920 1.00 20.93 AAAA ATOM 2868 C GLY 358 8.800 41.274 70.920 1.00 20.93 AAAA ATOM 2868 C GLY 358 8.800 41.274 70.920 1.00 20.93 AAAA ATOM 2868 C GLY 358 8.800 41.274 70.920 1.00 20.93 AAAA ATOM 2868 C GLY 358 8.800 41.274 70.920 1.00 20.93 AAAA ATOM 2889 C GLY 358 8.800 41.274 70.920 1.00 20.93 AAAA ATOM 2889 C GLY 368										
ATOM 2844 NEI TRP 355 19.765 46.991 72.997 1.00 56.07 AAAA ATOM 2846 C23 TRP 355 19.887 43.556 72.897 1.00 55.25 AAAA ATOM 2848 C TRP 355 19.887 43.356 72.405 1.00 50.68 AAAA ATOM 2848 C TRP 355 15.899 47.712 68.776 1.00 33.12 AAAA ATOM 2848 C TRP 355 15.996 48.705 68.550 1.00 29.41 AAAA ATOM 2850 N ARG 356 15.547 46.508 68.263 1.00 29.41 AAAA ATOM 2851 CA ARG 356 14.413 46.237 67.387 1.00 23.96 AAAA ATOM 2852 CB ARG 356 14.413 46.237 67.387 1.00 23.96 AAAA ATOM 2853 CG ARG 356 14.413 48.92 46.096 65.935 1.00 23.96 AAAA ATOM 2854 CD ARG 356 16.291 47.212 64.108 1.00 29.91 AAAA ATOM 2855 CZ ARG 356 16.891 47.212 64.108 1.00 29.92 AAAA ATOM 2855 CZ ARG 356 16.891 47.212 64.108 1.00 29.05 AAAA ATOM 2856 CZ ARG 356 16.891 47.212 64.108 1.00 29.05 AAAA ATOM 2857 NH1 ARG 356 18.209 47.616 62.066 1.00 24.73 AAAA ATOM 2858 NH2 ARG 356 18.209 47.616 62.066 1.00 22.15 AAAA ATOM 2858 NH2 ARG 356 18.209 47.616 62.066 1.00 22.15 AAAA ATOM 2858 NH2 ARG 356 18.209 47.616 62.066 1.00 22.15 AAAA ATOM 2858 NH2 ARG 356 18.209 47.616 62.066 1.00 22.15 AAAA ATOM 2860 O ARG 356 13.785 43.925 67.189 1.00 22.59 AAAA ATOM 2861 N GLY 357 12.631 43.805 69.657 1.00 22.59 AAAA ATOM 2862 CA GLY 357 12.631 43.805 69.657 1.00 22.99 AAAA ATOM 2868 O GLY 357 11.138 43.691 69.055 1.00 22.99 AAAA ATOM 2868 O GLY 357 11.538 44.994 69.055 1.00 22.95 AAAA ATOM 2868 O GLY 357 12.631 43.805 69.657 1.00 22.95 AAAA ATOM 2868 O GLY 357 12.631 43.805 69.657 1.00 22.95 AAAA ATOM 2868 O GLY 358 9.184 42.956 70.056 1.00 22.95 AAAA ATOM 2868 O GLY 357 12.631 43.805 69.657 1.00 22.95 AAAA ATOM 2868 O GLY 357 12.631 43.805 69.657 1.00 22.95 AAAA ATOM 2868 O GLY 358 9.184 42.957 70.188 1.00 22.97 AAAA ATOM 2868 O GLY 358 9.184 42.977 70.20 1.00 22.97 AAAA ATOM 2868 O GLY 358 9.569 9.507 1.00 22.97 AAAA ATOM 2868 O GLY 358 9.569 9.507 1.00 22.97 AAAA ATOM 2870 C G GLU 359 7.601 40.747 71.50 1.00 22.97 AAAA ATOM 2871 C G GLU 359 7.804 42.797 70.20 1.00 22.97 AAAA ATOM 2878 N VAL 360 9.305 9.307 71.366 1.00 22.97 AAAA ATOM 2878 O VAL 360 9.305 9.307 71.366 1.00 22.										
ATOM 2845 C22 TRP 355	ATOM	2843	CD	LTRP	355	18.723	47.746	72.507	1.00 56.10	AAAA
ATOM 2845 CZ2 TRP 355 20.340 44.552 72.897 1.00 55.25 AAAA ATOM 2846 CZ3 TRP 355 18.696 43.267 71.674 1.00 50.74 AAAA ATOM 2848 C TRP 355 19.887 43.356 72.405 1.00 50.68 AAAA ATOM 2848 C TRP 355 15.789 47.712 68.776 1.00 33.12 AAAA ATOM 2849 O TRP 355 15.789 47.712 68.776 1.00 23.94 AAAA ATOM 2850 N ARG 356 15.547 46.508 68.253 1.00 23.90 AAAA ATOM 2851 CA ARG 356 14.492 46.996 65.955 1.00 23.96 AAAA ATOM 2852 CB ARG 356 14.892 46.096 65.955 1.00 22.66 AAAA ATOM 2853 CG ARG 356 15.547 47.825 65.955 1.00 22.66 AAAA ATOM 2854 CD ARG 356 16.291 47.212 64.108 1.00 28.92 AAAA ATOM 2855 NF ARG 356 16.833 48.503 63.666 1.00 24.73 AAAA ATOM 2858 NFL ARG 356 17.7133 48.668 62.724 1.00 23.57 AAAA ATOM 2858 NFL ARG 356 18.153 49.891 62.724 1.00 22.55 AAAA ATOM 2858 NFL ARG 356 18.153 49.891 62.724 1.00 22.55 AAAA ATOM 2858 NFL ARG 356 18.153 49.891 62.064 1.00 22.55 AAAA ATOM 2858 NFL ARG 356 18.153 49.891 62.048 10.00 22.55 AAAA ATOM 2858 NFL ARG 356 13.781 44.944 67.878 1.00 22.55 AAAA ATOM 2858 NFL ARG 356 13.781 44.944 67.878 1.00 22.55 AAAA ATOM 2860 C ARG 356 33.781 44.944 67.878 1.00 22.55 AAAA ATOM 2861 N ARG 356 33.781 44.944 67.878 1.00 22.55 AAAA ATOM 2863 C CLLY 357 12.231 44.993 69.085 1.00 22.25 AAAA ATOM 2865 N SILY 358	ATOM	2844	NEI	LTRP	355	19.765	46.991	72.997	1.00 56.07	AAAA
ATOM 2846 C23 TRP 355 18.696 43.267 71.674 1.00 50.74 AAAA ATOM 2848 C TRP 355 19.887 43.356 72.405 1.00 50.68 AAAA ATOM 2848 C TRP 355 15.589 47.712 68.776 1.00 33.12 AAAA ATOM 2850 N ARG 356 15.547 46.508 68.263 1.00 23.90 AAAA ATOM 2851 CA ARG 356 14.813 46.237 67.387 1.00 23.90 AAAA ATOM 2852 CB ARG 356 14.892 46.096 65.935 1.00 22.66 AAAA ATOM 2852 CB ARG 356 14.892 46.096 65.935 1.00 22.66 AAAA ATOM 2853 CC ARG 356 16.891 47.385 65.393 1.00 22.66 AAAA ATOM 2854 CD ARG 356 16.891 47.212 64.108 1.00 23.96 AAAA ATOM 2855 NE ARG 356 16.891 47.212 64.108 1.00 28.92 AAAA ATOM 2855 NE ARG 356 16.891 47.212 64.108 1.00 23.96 AAAA ATOM 2855 NE ARG 356 16.891 47.212 64.108 1.00 23.57 AAAA ATOM 2855 NE ARG 356 16.891 47.212 64.108 1.00 22.56 AAAA ATOM 2856 C2 ARG 356 17.733 48.668 62.724 1.00 23.57 AAAA ATOM 2857 NH1 ARG 356 18.153 48.503 63.666 1.00 22.15 AAAA ATOM 2858 NH2 ARG 356 18.153 48.891 62.724 1.00 22.55 AAAA ATOM 2859 C ARG 356 13.781 44.944 67.878 1.00 22.55 AAAA ATOM 2850 C ARG 356 13.781 44.944 67.878 1.00 22.55 AAAA ATOM 2861 N GLY 357 12.631 44.993 69.085 1.00 22.25 AAAA ATOM 2862 C AGLY 357 12.631 44.993 69.085 1.00 22.25 AAAA ATOM 2863 C GLY 357 13.231 44.993 69.085 1.00 22.25 AAAA ATOM 2866 C AGLY 357 10.536 44.330 66.69 1.00 22.25 AAAA ATOM 2867 C GLY 358 8.800 41.274 70.920 1.00 23.91 AAAA ATOM 2868 N GLY 358 8.800 41.274 70.920 1.00 28.92 AAAA ATOM 2868 N GLY 358 8.800 41.274 70.920 1.00 30.03 AAAA ATOM 2869 N GLU 359 7.0536 8.800 41.274 70.920 1.00 30.03 AAAA ATOM 2869 N GLU 359 7.601 40.747 70.715 1.00 28.92 AAAA ATOM 2868 C GLY 358 8.800 41.274 70.920 1.00 30.03 AAAA ATOM 2869 N GLU 359 7.601 40.747 70.715 1.00 22.25 AAAA ATOM 2869 N GLU 359 7.601 40.747 70.715 1.00 22.99 AAAA ATOM 2869 N GLU 359 7.601 40.747 70.715 1.00 24.03 AAAA ATOM 2869 N GLU 359 7.601 40.747 70.715 1.00 24.03 AAAA ATOM 2877 C G GLU 359 8.918 8.918 8.919 1.00 22.94 AAAA ATOM 2878 C GLU 359 8.918 8.918 8.919 1.00 22.94 AAAA ATOM 2878 C GLU 359 8.918 8.918 8.919 1.00 22.91 AAAA ATOM 2888 C G ARG 361 5.949 1.00 22.91 AAAA										
ATOM 2847 CH2 TRP 355	_									
ATOM 2849 C TRP 355 15.789 47.712 68.776 1.00 33.12 AAAA ATOM 2850 N ARG 356 15.596 48.705 68.505 1.00 29.41 ATOM 2851 CA ARG 356 14.413 46.237 67.387 1.00 23.90 AAAA ATOM 2852 CB ARG 356 14.413 46.237 67.387 1.00 23.90 AAAA ATOM 2852 CB ARG 356 14.413 46.237 67.387 1.00 23.90 AAAA ATOM 2852 CB ARG 356 14.413 46.237 67.387 1.00 22.66 AAAA ATOM 2854 CD ARG 356 15.595 47.385 65.393 1.00 22.66 AAAA ATOM 2855 NC ARG 356 15.595 47.385 65.393 1.00 22.66 AAAA ATOM 2855 NC ARG 356 16.833 48.503 61.686 1.00 28.22 AAAA ATOM 2858 NH1 ARG 356 16.833 48.503 61.686 1.00 22.15 AAAA ATOM 2858 NH2 ARG 356 18.153 49.891 62.418 1.00 22.69 AAAA ATOM 2858 NH2 ARG 356 18.153 49.891 62.418 1.00 22.69 AAAA ATOM 2858 NH2 ARG 356 13.781 44.944 67.878 1.00 22.89 AAAA ATOM 2850 O ARG 356 13.781 44.944 67.878 1.00 22.89 AAAA ATOM 2861 N GLY 357 12.631 43.805 69.657 1.00 23.91 AAAA ATOM 2862 CA GLY 357 12.631 43.805 69.657 1.00 22.59 AAAA ATOM 2863 C GLY 357 12.631 43.805 69.657 1.00 26.90 AAAA ATOM 2864 O GLY 357 10.536 44.330 69.657 1.00 26.90 AAAA ATOM 2865 N GLY 358 10.544 42.797 70.265 1.00 28.22 AAAA ATOM 2866 CA GLY 358 8.800 41.274 70.920 1.00 30.03 AAAA ATOM 2867 C GLY 358 8.800 42.797 70.265 1.00 28.22 AAAA ATOM 2867 C GLY 358 8.800 42.797 70.265 1.00 28.22 AAAA ATOM 2868 O GLY 358 9.104 42.551 70.188 1.00 23.91 AAAAA ATOM 2867 C GLY 358 8.800 41.274 70.920 1.00 30.03 AAAA ATOM 2867 C GLU 359 7.601 40.747 70.715 1.00 26.90 AAAAA ATOM 2868 O GLY 358 9.104 42.551 70.188 1.00 29.91 AAAAA ATOM 2867 C GLU 359 7.601 40.747 70.715 1.00 26.92 AAAA ATOM 2868 O GLY 358 9.104 42.551 70.188 1.00 29.92 AAAA ATOM 2867 C GLU 359 7.601 40.747 70.715 1.00 29.52 AAAAA ATOM 2868 O GLY 358 9.606 40.757 71.663 1.00 24.03 AAAA ATOM 2873 CD GLU 359 7.601 40.747 70.715 1.00 29.52 AAAAA ATOM 2873 CD GLU 359 7.601 40.747 70.715 1.00 29.52 AAAAA ATOM 2873 CD GLU 359 7.601 40.747 70.715 1.00 29.52 AAAAA ATOM 2874 OEI GLU 359 9.609 39.372 71.375 1.00 29.52 AAAAA ATOM 2878 N AAAA ATOM 2879 CA VAL 360 9.783 31.778 89.70 29.91 1.00 30.03 AAAAA ATOM 2889 CD ARG 3										
ATOM 2849 O TRP 355 15.096 48.705 68.550 1.00 29.41 AAAA ATOM 2851 CA ARG 356 15.547 46.598 68.263 1.00 23.96 AAAA ATOM 2851 CCA ARG 356 14.913 46.237 67.387 1.00 23.96 AAAA ATOM 2852 CB ARG 356 14.992 46.096 65.935 1.00 29.06 AAAA ATOM 2853 CG ARG 356 15.505 47.385 65.393 1.00 29.06 AAAA ATOM 2854 CD ARG 356 16.291 47.385 65.393 1.00 29.06 AAAA ATOM 2855 NE ARG 356 16.893 47.212 64.108 1.00 28.92 AAAA ATOM 2856 CZ ARG 356 17.733 48.668 62.724 1.00 23.57 AAAA ATOM 2857 NNII ARG 356 18.209 47.616 62.066 1.00 22.15 ATOM 2858 NNI2 ARG 356 18.209 47.616 62.066 1.00 22.59 AAAA ATOM 2858 NNI2 ARG 356 13.781 44.944 67.878 1.00 22.69 AAAA ATOM 2850 C ARG 356 13.785 43.925 67.189 1.00 22.69 AAAA ATOM 2860 O ARG 356 13.785 43.925 67.189 1.00 22.25 AAAA ATOM 2860 O ARG 356 13.785 43.805 69.657 1.00 26.72 AAAA ATOM 2860 C ARG 356 13.785 43.805 69.657 1.00 26.72 AAAA ATOM 2864 O GLY 357 11.138 43.671 69.465 1.00 29.87 ATOM 2866 C ARG 358 10.536 44.330 68.619 1.00 29.87 AATOM 2866 C ARG 358 8 10.534 42.797 70.265 1.00 26.72 AAAA ATOM 2866 C ARG 358 9.118 42.561 70.188 1.00 24.03 AATOM 2866 C ARG 358 8 8.800 41.274 70.920 1.00 30.96 AAAA ATOM 2867 C G LY 358 9.118 42.561 70.188 1.00 24.03 AATOM 2868 O GLY 358 9.626 40.757 71.663 1.00 24.03 AATOM 2868 O GLY 358 9.626 40.757 71.663 1.00 24.03 AATOM 2868 O GLY 359 7.218 39.509 71.366 1.00 24.03 AATOM 2868 O GLY 359 7.218 39.509 71.366 1.00 24.03 AATOM 2868 O GLY 359 7.218 39.509 71.366 1.00 24.03 AATOM 2868 O GLY 359 7.218 39.509 71.366 1.00 24.03 AATOM 2868 O GLY 359 7.218 39.509 71.366 1.00 24.03 AATOM 2868 O GLY 359 7.218 39.509 71.366 1.00 24.03 AATOM 2868 O GLY 359 7.218 39.509 71.366 1.00 24.03 AATOM 2870 CA GLU 359 7.218 39.509 71.375 1.00 25.51 AAAA ATOM 2871 CR GLU 359 7.218 39.509 71.366 1.00 24.03 AATOM 2878 N ARG 361 7.944 37.208 69.657 1.00 25.51 AAAA ATOM 2878 N ARG 361 7.944 37.208 69.125 1.00 2.128 AATOM 2878 N ARG 361 7.944 37.208 69.125 1.00 2.128 AATOM 2878 N ARG 361 5.947 34.337 66.221 1.00 2.128 AATOM 2880 CD ARG 361 5.948 34.389 66.772 1.00 2.128 AATOM 2880	ATOM	2847	CHZ	TRP	355	19.887	43.356	72.405	1.00 50.68	AAAA
ATOM 2849 O TRP 355 15.096 48.705 68.550 1.00 29.41 AAAA ATOM 2851 CA ANG 356 15.547 46.508 68.263 1.00 23.96 AAAA ATOM 2851 CA ANG 356 14.992 46.096 68.263 1.00 23.96 AAAA ATOM 2852 CB ANG 356 14.992 46.096 65.935 1.00 22.66 AAAA ATOM 2853 CG ANG 356 15.505 47.385 65.393 1.00 22.06 AAAA ATOM 2855 NE ANG 356 16.833 48.503 64.108 1.00 23.96 AAAA ATOM 2855 NE ANG 356 16.833 48.503 64.108 1.00 24.90 AAAA ATOM 2855 NE ANG 356 16.833 48.503 64.108 1.00 24.90 AAAA ATOM 2855 NE ANG 356 16.833 48.503 64.108 1.00 24.90 AAAA ATOM 2857 NNII ANG 356 18.209 47.616 62.066 1.00 22.57 AAAA ATOM 2858 NE ANG 356 18.209 47.616 62.066 1.00 22.59 AAAA ATOM 2858 NE ANG 356 13.781 44.944 67.878 1.00 22.69 AAAA ATOM 2850 CA GUY 357 13.231 44.993 69.085 1.00 29.87 AAAA ATOM 2860 O ANG 356 13.785 43.925 67.189 1.00 22.25 AAAA ATOM 2860 O ANG 356 13.785 43.805 69.657 1.00 26.72 AAAA ATOM 2864 O GUY 357 11.138 43.671 69.465 1.00 29.87 AAAA ATOM 2866 CA GUY 357 11.138 43.671 69.465 1.00 29.87 AAAA ATOM 2866 CA GUY 357 10.536 44.330 68.619 1.00 29.87 AAAA ATOM 2866 CY ANG 356 88.800 41.274 70.920 1.00 30.96 AAAA ATOM 2868 O GUY 358 9.118 42.561 70.188 1.00 30.96 AAAA ATOM 2868 O GUY 358 9.118 42.561 70.188 1.00 30.96 AAAA ATOM 2868 O GUY 358 9.626 40.757 71.663 1.00 24.03 AAAA ATOM 2868 O GUY 358 9.626 40.757 71.663 1.00 24.03 AAAA ATOM 2868 O GUY 358 9.626 40.757 71.663 1.00 24.03 AAAA ATOM 2868 O GUY 359 7.601 40.747 70.715 1.00 28.34 AAAA ATOM 2870 CA GUU 359 7.614 40.757 71.663 1.00 24.03 AAAA ATOM 2878 N VAL 360 9.783 33.917 70.912 1.00 30.96 AAAA ATOM 2878 N VAL 360 9.783 33.917 70.912 1.00 30.93 AAAA ATOM 2878 N VAL 360 9.783 33.917 70.912 1.00 30.93 AAAA ATOM 2878 N VAL 360 9.783 33.917 70.912 1.00 30.93 AAAA ATOM 2878 N VAL 360 9.783 33.917 70.912 1.00 30.95 AAAA ATOM 2878 N VAL 360 9.783 33.917 70.912 1.00 22.94 AAAA ATOM 2878 N VAL 360 9.783 33.917 70.912 1.00 22.94 AAAA ATOM 2878 N VAL 360 9.783 33.917 70.912 1.00 22.94 AAAA ATOM 2880 CB VAL 360 9.300 35.188 71.621 1.00 22.91 AAAA ATOM 2881 CG VAL 360 9.300 35.188 71.621 1.00 22	MOTA	2848	С	TRP	355	15.789	47.712	68.776	1.00 33.12	AAAA
ATOM 2850 N ARG 356 14.413 46.237 67.387 1.00 23.90 AAAA ATOM 2851 CA ARG 356 14.413 46.237 67.387 1.00 23.90 AAAA ATOM 2851 CC ARG 356 14.4192 46.096 65.935 1.00 22.66 AAAA ATOM 2853 CG ARG 356 15.505 47.385 65.393 1.00 22.66 AAAA ATOM 2855 NE ARG 356 16.291 47.212 64.108 1.00 28.92 AAAA ATOM 2856 CZ ARG 356 16.291 47.212 64.108 1.00 28.92 AAAA ATOM 2857 NNH ARG 356 17.733 48.668 62.724 1.00 23.57 AAAA ATOM 2858 NN2 ARG 356 18.153 49.891 62.418 1.00 22.15 AAAA ATOM 2858 NN2 ARG 356 18.153 49.891 62.418 1.00 22.69 AAAA ATOM 2859 CZ ARG 356 18.153 49.891 62.418 1.00 22.69 AAAA ATOM 2859 CZ ARG 356 13.781 44.944 67.878 1.00 22.89 AAAA ATOM 2861 N GLY 357 12.631 43.805 69.657 1.00 23.91 AAAA ATOM 2863 CZ GLY 357 12.631 43.805 69.657 1.00 26.90 AAAA ATOM 2863 CZ GLY 357 12.631 43.805 69.657 1.00 26.90 AAAA ATOM 2863 CZ GLY 357 10.536 44.330 68.619 1.00 22.89 AAAA ATOM 2865 N GLY 358 10.544 42.797 70.265 1.00 26.90 AAAA ATOM 2869 N GLY 358 9.114 42.561 70.188 1.00 10.0 28.22 AAAA ATOM 2867 N GLY 358 9.114 42.561 70.188 1.00 10.0 28.22 AAAA ATOM 2867 C GLY 358 9.114 42.561 70.188 1.00 10.0 28.22 AAAA ATOM 2867 C GLY 358 9.118 42.561 70.188 1.00 10.0 28.22 AAAA ATOM 2867 C GLY 358 9.118 42.561 70.188 1.00 10.0 28.22 AAAA ATOM 2867 C GLY 358 9.118 42.561 70.188 1.00 10.0 3 AAAA ATOM 2867 C GLY 359 9.10 359 7.601 40.747 70.715 1.00 26.23 AAAA ATOM 2867 C GLY 359 9.10 359 7.601 40.747 70.715 1.00 26.23 AAAA ATOM 2867 C GLU 359 7.218 39.377 71.663 1.00 24.03 AAAA ATOM 2867 C GLU 359 9.00 44.91 70.715 1.00 24.37 AAAA ATOM 2870 C GLU 359 9.00 44.91 70.715 1.00 24.37 AAAA ATOM 2877 C G GLU 359 9.00 44.91 70.715 1.00 24.37 AAAA ATOM 2877 C GLU 359 9.30 377 71.663 1.00 24.03 AAAA ATOM 2878 N VAL 360 9.30 3.3172 40.250 71.375 1.00 32.52 AAAA ATOM 2878 N VAL 360 9.30 35.188 71.021 1.00 24.37 AAAA ATOM 2878 N VAL 360 9.30 35.188 71.021 1.00 24.37 AAAA ATOM 2878 N VAL 360 9.30 35.188 71.021 1.00 25.79 AAAA ATOM 2886 C AARG 361 5.949 34.317 66.224 1.00 22.91 AAAA ATOM 2887 C BAG 361 5.949 34.317 66.224 1.00 22.91 AAAA ATOM 2889 C B							48 705			
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ATOM 2874 OE1 GLU 359 2.924 39.125 72.151 1.00 42.92 AAAA ATOM 2875 OE2 GLU 359 7.804 38.323 70.628 1.00 27.35 AAAA ATOM 2877 O GLU 359 8.138 38.415 69.449 1.00 22.94 AAAA ATOM 2878 N VAL 360 7.944 37.208 71.325 1.00 19.68 AAAA ATOM 2879 CA VAL 360 8.441 36.017 70.672 1.00 19.68 AAAA ATOM 2879 CA VAL 360 9.300 35.188 71.621 1.00 26.71 AAAA ATOM 2880 CB VAL 360 9.300 35.188 71.621 1.00 26.71 AAAA ATOM 2881 CG1 VAL 360 9.783 33.917 70.912 1.00 20.64 AAAA ATOM 2882 CG2 VAL 360 10.886 36.038 72.113 1.00 25.79 AAAA ATOM 2883 C VAL 360 10.886 36.038 72.113 1.00 25.79 AAAA ATOM 2884 O VAL 360 7.228 35.202 70.197 1.00 25.51 AAAA ATOM 2884 O VAL 360 6.442 34.700 71.01 1.00 19.75 AAAA ATOM 2885 N ARG 361 7.065 35.094 68.85 1.00 19.75 AAAA ATOM 2887 CB ARG 361 5.947 34.337 68.3C 1.00 22.01 AAAA ATOM 2888 CG ARG 361 5.947 34.337 68.3C 1.00 22.01 AAAA ATOM 2888 CG ARG 361 5.947 34.337 68.3C 1.00 22.01 AAAA ATOM 2889 CD ARG 361 5.948 34.389 66.772 1.00 19.31 AAAA ATOM 2889 CD ARG 361 5.735 35.730 64.723 1.00 30.86 AAAA ATOM 2890 NE ARG 361 5.735 35.730 64.723 1.00 30.86 AAAA ATOM 2891 CZ ARG 361 7.616 36.275 63.242 1.00 22.89 AAAA ATOM 2892 NH1 ARG 361 8.861 36.704 63.081 1.00 22.89 AAAA ATOM 2897 CA ARG 361 5.897 32.879 68.714 1.00 22.89 AAAA ATOM 2897 CA ARG 361 5.897 32.879 68.714 1.00 22.89 AAAA ATOM 2897 CA ARG 361 5.897 32.879 68.714 1.00 22.89 AAAA ATOM 2897 CA ARG 361 5.897 32.879 68.714 1.00 22.89 AAAA ATOM 2897 CA ARG 361 5.897 32.879 68.714 1.00 22.179 AAAA ATOM 2897 CA ARS 362 4.681 32.338 68.763 1.00 24.89 AAAA ATOM 2897 CA ARS 362 4.681 32.338 68.763 1.00 24.89 AAAA ATOM 2898 CB ARG 361 5.897 32.879 68.714 1.00 22.991 AAAA ATOM 2899 CG ARS 362 2.981 30.570 69.070 1.00 22.991 AAAA ATOM 2899 CG ARS 362 2.981 30.570 69.070 1.00 22.991 AAAA ATOM 2899 CG ARS 362 2.981 30.570 69.070 1.00 22.991 AAAA ATOM 2890 CC ARS 362 2.981 30.570 69.070 1.00 22.991 AAAA ATOM 2890 CC ARS 362 2.981 30.570 69.070 1.00 22.991 AAAA ATOM 2890 CC ARS 362 2.2961 30.570 69.070 1.00 22.991 AAAA ATOM 2900 CD ARS 362 2.2961 30.570 69.070 1.00	ATOM	2872	CG	GLU	359	4.981	40:327	72.299	1.00 45.44	AAAA
ATOM 2874 OE1 GLU 359 2.924 39.125 72.151 1.00 42.92 AAAA ATOM 2875 OE2 GLU 359 7.804 38.323 70.628 1.00 27.35 AAAA ATOM 2877 O GLU 359 8.138 38.415 69.449 1.00 22.94 AAAA ATOM 2878 N VAL 360 7.944 37.208 71.325 1.00 19.68 AAAA ATOM 2879 CA VAL 360 8.441 36.017 70.672 1.00 19.68 AAAA ATOM 2879 CA VAL 360 9.300 35.188 71.621 1.00 26.71 AAAA ATOM 2880 CB VAL 360 9.300 35.188 71.621 1.00 26.71 AAAA ATOM 2881 CG1 VAL 360 9.783 33.917 70.912 1.00 20.64 AAAA ATOM 2882 CG2 VAL 360 10.886 36.038 72.113 1.00 25.79 AAAA ATOM 2883 C VAL 360 10.886 36.038 72.113 1.00 25.79 AAAA ATOM 2884 O VAL 360 7.228 35.202 70.197 1.00 25.51 AAAA ATOM 2884 O VAL 360 6.442 34.700 71.01 1.00 19.75 AAAA ATOM 2885 N ARG 361 7.065 35.094 68.85 1.00 19.75 AAAA ATOM 2887 CB ARG 361 5.947 34.337 68.3C 1.00 22.01 AAAA ATOM 2888 CG ARG 361 5.947 34.337 68.3C 1.00 22.01 AAAA ATOM 2888 CG ARG 361 5.947 34.337 68.3C 1.00 22.01 AAAA ATOM 2889 CD ARG 361 5.948 34.389 66.772 1.00 19.31 AAAA ATOM 2889 CD ARG 361 5.735 35.730 64.723 1.00 30.86 AAAA ATOM 2890 NE ARG 361 5.735 35.730 64.723 1.00 30.86 AAAA ATOM 2891 CZ ARG 361 7.616 36.275 63.242 1.00 22.89 AAAA ATOM 2892 NH1 ARG 361 8.861 36.704 63.081 1.00 22.89 AAAA ATOM 2897 CA ARG 361 5.897 32.879 68.714 1.00 22.89 AAAA ATOM 2897 CA ARG 361 5.897 32.879 68.714 1.00 22.89 AAAA ATOM 2897 CA ARG 361 5.897 32.879 68.714 1.00 22.89 AAAA ATOM 2897 CA ARG 361 5.897 32.879 68.714 1.00 22.89 AAAA ATOM 2897 CA ARG 361 5.897 32.879 68.714 1.00 22.179 AAAA ATOM 2897 CA ARS 362 4.681 32.338 68.763 1.00 24.89 AAAA ATOM 2897 CA ARS 362 4.681 32.338 68.763 1.00 24.89 AAAA ATOM 2898 CB ARG 361 5.897 32.879 68.714 1.00 22.991 AAAA ATOM 2899 CG ARS 362 2.981 30.570 69.070 1.00 22.991 AAAA ATOM 2899 CG ARS 362 2.981 30.570 69.070 1.00 22.991 AAAA ATOM 2899 CG ARS 362 2.981 30.570 69.070 1.00 22.991 AAAA ATOM 2890 CC ARS 362 2.981 30.570 69.070 1.00 22.991 AAAA ATOM 2890 CC ARS 362 2.981 30.570 69.070 1.00 22.991 AAAA ATOM 2890 CC ARS 362 2.2961 30.570 69.070 1.00 22.991 AAAA ATOM 2900 CD ARS 362 2.2961 30.570 69.070 1.00	ATOM	2873	CD	GLU	359	3.472	40.250	72.132	1.00 50.43	AAAA
ATOM 2875 OE2 GLU 359 7.804 38.323 70.628 1.00 40.72 AAAA ATOM 2876 C GLU 359 7.804 38.323 70.628 1.00 27.35 AAAA ATOM 2877 O GLU 359 8.138 38.415 69.449 1.00 22.94 AAAA ATOM 2878 N VAL 360 7.944 37.208 71.325 1.00 19.68 AAAA ATOM 2879 CA VAL 360 8.441 36.017 70.672 1.00 21.28 AAAA ATOM 2880 CB VAL 360 9.300 35.188 71.621 1.00 26.71 AAAA ATOM 2881 CG1 VAL 360 9.300 35.188 71.621 1.00 26.71 AAAA ATOM 2882 CG2 VAL 360 9.783 33.917 70.912 1.00 20.64 AAAA ATOM 2883 C VAL 360 9.783 33.917 70.912 1.00 25.79 AAAA ATOM 2883 C VAL 360 7.228 35.202 70.197 1.00 25.79 AAAA ATOM 2884 O VAL 360 6.442 34.700 71.01 1.00 19.75 AAAA ATOM 2885 N ARG 361 7.065 35.094 68.813 1.00 18.48 AAAA ATOM 2887 CB ARG 361 5.947 34.337 68.30 1.00 18.48 AAAA ATOM 2887 CB ARG 361 5.947 34.337 68.30 1.00 22.01 AAAA ATOM 2888 CG ARG 361 5.947 34.337 68.30 1.00 22.01 AAAA ATOM 2888 CG ARG 361 5.446 35.671 66.204 1.00 30.86 AAAA ATOM 2889 CD ARG 361 5.446 35.671 66.204 1.00 30.86 AAAA ATOM 2889 CD ARG 361 5.446 35.671 66.204 1.00 30.73 AAAA ATOM 2891 CZ ARG 361 7.616 36.275 63.242 1.00 22.89 AAAA ATOM 2893 NH1 ARG 361 6.851 36.006 62.186 1.00 19.02 AAAA ATOM 2893 NH2 ARG 361 7.616 36.275 63.242 1.00 22.89 AAAA ATOM 2893 NH2 ARG 361 5.897 32.879 68.714 1.00 23.47 AAAA ATOM 2893 NH2 ARG 361 5.897 32.879 68.714 1.00 23.47 AAAA ATOM 2893 NH2 ARG 361 5.897 32.879 68.714 1.00 23.47 AAAA ATOM 2893 NH2 ARG 361 5.897 32.879 68.714 1.00 23.47 AAAA ATOM 2893 C ARG 361 5.897 32.879 68.714 1.00 22.91 AAAA ATOM 2897 C ARG 361 5.897 32.879 68.714 1.00 23.47 AAAA ATOM 2897 C ARG 361 5.897 32.879 68.714 1.00 22.91 AAAA ATOM 2898 CB LYS 362 2.961 30.570 69.070 1.00 22.91 AAAA ATOM 2899 CG LYS 362 2.961 30.570 69.070 1.00 22.91 AAAA ATOM 2899 CG LYS 362 2.961 30.570 69.070 1.00 22.91 AAAA ATOM 2899 CG LYS 362 2.996 30.014 68.202 1.00 16.77 AAAA ATOM 2900 CD LYS 362 2.397 34.683 68.711 1.00 22.99 AAAA ATOM 2900 CD LYS 362 2.397 34.683 68.711 1.00 22.99 AAAA ATOM 2900 CD LYS 362 2.397 34.683 68.711 1.00 22.99 AAAA ATOM 2900 CD LYS 362 2.307 34.683 68.711 1.00 22.99 AAAA										
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ATOM 2879 CA VAL 360 8.441 36.017 70.672 1.00 21.28 AAAA ATOM 2880 CB VAL 360 9.300 35.188 71.621 1.00 26.71 AAAA ATOM 2881 CG1 VAL 360 9.783 33.917 70.912 1.00 20.64 AAAA ATOM 2882 CG2 VAL 360 10.486 36.038 72.113 1.00 25.79 AAAA ATOM 2883 C VAL 360 7.228 35.202 70.197 1.00 25.51 AAAA ATOM 2884 O VAL 360 6.442 34.700 71.01 1.00 19.75 AAAA ATOM 2885 N ARG 361 7.065 35.094 68.8.3 1.00 19.75 AAAA ATOM 2886 CA ARG 361 5.947 34.337 68.3C 1.00 22.01 AAAA ATOM 2887 CB ARG 361 5.988 34.389 66.772 1.00 19.31 AAAA ATOM 2888 CG ARG 361 5.988 34.389 66.772 1.00 19.31 AAAA ATOM 2889 CD ARG 361 5.735 35.730 64.723 1.00 30.86 AAAA ATOM 2889 CD ARG 361 7.111 36.148 64.460 1.00 30.73 AAAA ATOM 2890 NE ARG 361 7.111 36.148 64.460 1.00 30.73 AAAA ATOM 2892 NH1 ARG 361 7.616 36.275 63.242 1.00 22.89 AAAA ATOM 2893 NH2 ARG 361 6.851 36.006 62.186 1.00 19.02 AAAA ATOM 2893 NH2 ARG 361 8.861 36.704 63.081 1.00 23.47 AAAA ATOM 2894 C ARG 361 5.897 32.879 68.714 1.00 23.47 AAAA ATOM 2895 O ARG 361 6.851 36.006 62.186 1.00 19.02 AAAA ATOM 2897 CA LYS 362 4.681 32.338 68.763 1.00 24.89 AAAA ATOM 2897 CA LYS 362 4.681 32.338 68.763 1.00 24.89 AAAA ATOM 2898 CB LYS 362 4.479 30.938 69.125 1.00 22.91 AAAA ATOM 2898 CB LYS 362 2.981 30.570 69.070 1.00 22.91 AAAA ATOM 2899 CG LYS 362 2.981 30.570 69.070 1.00 22.91 AAAA ATOM 2899 CG LYS 362 2.981 30.570 69.070 1.00 22.91 AAAA ATOM 2899 CG LYS 362 2.981 30.570 69.070 1.00 22.91 AAAA ATOM 2899 CG LYS 362 2.981 30.570 69.070 1.00 22.91 AAAA ATOM 2899 CG LYS 362 2.981 30.570 69.070 1.00 22.91 AAAA ATOM 2900 CD LYS 362 2.290 32.715 70.157 1.00 57.51 AAAA ATOM 2900 CD LYS 362 2.290 32.715 70.157 1.00 50.87 AAAA ATOM 2900 CD LYS 362 2.290 32.715 70.157 1.00 50.87 AAAA ATOM 2900 CD LYS 362 2.290 32.715 70.157 1.00 50.87 AAAA ATOM 2900 CD LYS 362 2.290 32.715 70.157 1.00 50.87 AAAA ATOM 2900 CD LYS 362 2.307 34.683 68.711 1.00 22.99 AAAA ATOM 2900 CD LYS 362 2.307 34.683 68.711 1.00 22.99 AAAA	ATOM	2878	N	VAL	360	7.944	37.208	71.325	1.00 19.68	AAAA
ATOM 2880 CB VAL 360 9.300 35.188 71.621 1.00 26.71 AAAA ATOM 2881 CG1 VAL 360 9.783 33.917 70.912 1.00 20.64 AAAA ATOM 2882 CG2 VAL 360 10.486 36.038 72.113 1.00 25.79 AAAA ATOM 2883 C VAL 360 7.228 35.202 70.197 1.00 25.51 AAAA ATOM 2884 O VAL 360 6.442 34.700 71.01 1.00 19.75 AAAA ATOM 2885 N ARG 361 7.065 35.094 68.87.5 1.00 18.48 AAAA ATOM 2886 CA ARG 361 5.947 34.337 68.3C 1.00 22.01 AAAA ATOM 2887 CB ARG 361 5.948 34.389 66.772 1.00 19.31 AAAA ATOM 2888 CG ARG 361 5.946 35.671 66.204 1.00 30.86 AAAA ATOM 2889 CD ARG 361 5.446 35.671 66.204 1.00 30.86 AAAA ATOM 2889 CD ARG 361 5.735 35.730 64.723 1.00 37.95 AAAA ATOM 2890 NE ARG 361 7.111 36.148 64.460 1.00 30.73 AAAA ATOM 2891 CZ ARG 361 7.616 36.275 63.242 1.00 22.89 AAAA ATOM 2893 NH2 ARG 361 6.851 36.006 62.186 1.00 19.02 AAAA ATOM 2893 NH2 ARG 361 8.861 36.704 63.081 1.00 23.47 AAAA ATOM 2894 C ARG 361 5.897 32.879 68.714 1.00 26.11 AAAA ATOM 2894 C ARG 361 5.897 32.879 68.714 1.00 23.47 AAAA ATOM 2895 O ARG 361 6.926 32.255 68.968 1.00 21.79 AAAA ATOM 2897 CA LYS 362 4.681 32.338 68.763 1.00 24.89 AAAA ATOM 2898 CB LYS 362 4.479 30.938 69.125 1.00 22.91 AAAA ATOM 2898 CB LYS 362 4.479 30.938 69.125 1.00 22.91 AAAA ATOM 2898 CB LYS 362 2.961 30.570 69.070 1.00 22.91 AAAA ATOM 2899 CG LYS 362 2.961 30.570 69.070 1.00 22.91 AAAA ATOM 2899 CG LYS 362 2.961 30.570 69.070 1.00 22.91 AAAA ATOM 2899 CG LYS 362 2.290 32.715 70.157 1.00 57.51 AAAA ATOM 2900 CD LYS 362 2.290 32.715 70.157 1.00 50.86 AAAA ATOM 2900 CD LYS 362 2.290 32.715 70.157 1.00 50.87 AAAA ATOM 2900 NZ LYS 362 2.290 32.715 70.157 1.00 50.87 AAAA ATOM 2900 NZ LYS 362 2.290 32.715 70.157 1.00 50.87 AAAA ATOM 2900 NZ LYS 362 2.290 32.715 70.157 1.00 50.87 AAAA ATOM 2900 NZ LYS 362 2.307 34.683 68.799 1.00 50.87 AAAA ATOM 2900 NZ LYS 362 2.307 34.683 68.711 1.00 22.99 AAAA ATOM 2900 NZ LYS 362 2.307 34.683 68.711 1.00 22.99 AAAA ATOM 2900 NZ LYS 362 2.307 34.683 68.711 1.00 22.99 AAAA	ATOM	2879	CA	VAL	360	8.441	36.017	70.672		
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ATOM 2885 N ARG 361 7.065 35.094 68.8.3 1.00 18.48 AAAA ATOM 2886 CA ARG 361 5.947 34.337 68.3C 1.00 22.01 AAAA ATOM 2887 CB ARG 361 5.988 34.389 66.772 1.00 19.31 AAAA ATOM 2888 CG ARG 361 5.446 35.671 66.204 1.00 30.86 AAAA ATOM 2889 CD ARG 361 7.111 36.148 64.460 1.00 37.95 AAAA ATOM 2891 CZ ARG 361 7.616 36.275 63.242 1.00 22.89 AAAA ATOM 2892 NH1 ARG 361 7.616 36.275 63.242 1.00 22.89 AAAA ATOM 2893 NH2 ARG 361 8.861 36.704 63.081 1.00 23.47 AAAA ATOM 2894 C ARG 361 8.861 36.704 63.081 1.00 23.47 AAAA ATOM 2895 O ARG 361 5.897 32.879 68.714 1.00 26.11 AAAA ATOM 2896 N LYS 362 4.681 32.338 68.763 1.00 24.89 AAAA ATOM 2897 CA LYS 362 4.479 30.938 69.125 1.00 28.63 AAAA ATOM 2898 CB LYS 362 2.981 30.570 69.070 1.00 22.91 AAAA ATOM 2899 CG LYS 362 2.145 31.200 70.168 1.00 50.86 AAAA ATOM 2900 CD LYS 362 2.290 32.715 70.157 1.00 57.51 AAAA ATOM 2901 CE LYS 362 1.923 33.278 68.791 1.00 22.99 AAAA ATOM 2902 NZ LYS 362 2.307 34.683 68.711 1.00 22.99 AAAA ATOM 2901 CE LYS 362 2.290 32.715 70.157 1.00 57.51 AAAA ATOM 2902 NZ LYS 362 2.307 34.683 68.711 1.00 22.99 AAAA ATOM 2902 NZ LYS 362 2.307 34.683 68.711 1.00 22.99 AAAA ATOM 2903 C LYS 362 2.307 34.683 68.711 1.00 22.99 AAAA ATOM 2903 C LYS 362 5.269 30.014 68.202 1.00 16.77 AAAA	MOTA	2884		VAI.	360	6.442	34.700	71.011	1.00 19.75	AAAA
ATOM 2886 CA ARG 361 5.947 34.337 68.3C 1.00 22.01 AAAA ATOM 2887 CB ARG 361 5.988 34.389 66.772 1.00 19.31 AAAA ATOM 2888 CG ARG 361 5.446 35.671 66.204 1.00 30.86 AAAA ATOM 2889 CD ARG 361 5.735 35.730 64.723 1.00 37.95 AAAA ATOM 2890 NE ARG 361 7.111 36.148 64.460 1.00 30.73 AAAA ATOM 2891 CZ ARG 361 7.616 36.275 63.242 1.00 22.89 AAAA ATOM 2892 NH1 ARG 361 6.851 36.006 62.186 1.00 19.02 AAAA ATOM 2893 NH2 ARG 361 8.861 36.704 63.081 1.00 23.47 AAAA ATOM 2894 C ARG 361 8.861 36.704 63.081 1.00 23.47 AAAA ATOM 2895 O ARG 361 5.897 32.879 68.714 1.00 26.11 AAAA ATOM 2896 N LYS 362 4.681 32.338 68.763 1.00 21.79 AAAA ATOM 2897 CA LYS 362 4.681 32.338 68.763 1.00 24.89 AAAA ATOM 2898 CB LYS 362 4.479 30.938 69.125 1.00 28.63 AAAA ATOM 2898 CB LYS 362 2.981 30.570 69.070 1.00 22.91 AAAA ATOM 2899 CG LYS 362 2.145 31.200 70.168 1.00 50.86 AAAAA ATOM 2900 CD LYS 362 2.290 32.715 70.157 1.00 57.51 AAAAA ATOM 2901 CE LYS 362 2.290 32.715 70.157 1.00 57.51 AAAAA ATOM 2902 NZ LYS 362 2.307 34.683 68.711 1.00 22.99 AAAAA ATOM 2902 NZ LYS 362 2.307 34.683 68.711 1.00 22.99 AAAAA ATOM 2903 C LYS 362 2.307 34.683 68.711 1.00 22.99 AAAAA ATOM 2903 C LYS 362 5.269 30.014 68.202 1.00 16.77 AAAAA ATOM 2903 C LYS 362 5.269 30.014 68.202 1.00 16.77										
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ATOM 2890 NE ARG 361 7.111 36.148 64.460 1.00 30.73 AAAA ATOM 2891 CZ ARG 361 7.616 36.275 63.242 1.00 22.89 AAAA ATOM 2892 NH1 ARG 361 6.851 36.006 62.186 1.00 19.02 AAAA ATOM 2894 C ARG 361 8.861 36.704 63.081 1.00 23.47 AAAA ATOM 2895 O ARG 361 5.897 32.879 68.714 1.00 26.11 AAAA ATOM 2895 O ARG 361 6.926 32.255 68.968 1.00 21.79 AAAA ATOM 2896 N LYS 362 4.681 32.338 68.763 1.00 24.89 AAAA ATOM 2897 CA LYS 362 4.479 30.938 69.125 1.00 28.63 AAAA ATOM 2898 CB LYS 362 2.961 30.570 69.070 1.00 22.91 AAAA ATOM 2899 CG LYS 362 2.961 30.570 69.070 1.00 22.91 AAAA ATOM 2900 CD LYS 362 2.145 31.200 70.168 1.00 50.86 AAAA ATOM 2901 CE LYS 362 2.290 32.715 70.157 1.00 57.51 AAAAA ATOM 2901 CE LYS 362 2.307 34.683 68.791 1.00 22.99 AAAA ATOM 2902 NZ LYS 362 2.307 34.683 68.791 1.00 22.99 AAAA ATOM 2902 NZ LYS 362 2.307 34.683 68.711 1.00 22.99 AAAA ATOM 2903 C LYS 362 5.269 30.014 68.202 1.00 16.77 AAAA	ATOM	2889	CD	ARG	361	5.735	35.730	64.723	1.00 37.95	AAAA
ATOM 2891 CZ ARG 361 7.616 36.275 63.242 1.00 22.89 AAAA ATOM 2892 NH1 ARG 361 6.851 36.006 62.186 1.00 19.02 AAAA ATOM 2894 C ARG 361 8.861 36.704 63.081 1.00 23.47 AAAA ATOM 2895 O ARG 361 5.897 32.879 68.714 1.00 26.11 AAAA ATOM 2895 O ARG 361 6.926 32.255 68.968 1.00 21.79 AAAA ATOM 2896 N LYS 362 4.681 32.338 68.763 1.00 24.89 AAAA ATOM 2897 CA LYS 362 4.479 30.938 69.125 1.00 28.63 AAAA ATOM 2898 CB LYS 362 2.961 30.570 69.070 1.00 22.91 AAAA ATOM 2899 CG LYS 362 2.145 31.200 70.168 1.00 50.86 AAAA ATOM 2900 CD LYS 362 2.290 32.715 70.157 1.00 57.51 AAAAA ATOM 2901 CE LYS 362 1.923 33.278 68.799 1.00 50.87 AAAA ATOM 2902 NZ LYS 362 2.307 34.683 68.791 1.00 22.99 AAAA ATOM 2902 NZ LYS 362 2.307 34.683 68.711 1.00 22.99 AAAA ATOM 2903 C LYS 362 5.269 30.014 68.202 1.00 16.77 AAAA										2222
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ATOM 2895 O ARG 361 6.926 32.255 68.968 1.00 21.79 AAAA ATOM 2896 N LYS 362 4.681 32.338 68.763 1.00 24.89 AAAA ATOM 2897 CA LYS 362 4.479 30.938 69.125 1.00 28.63 AAAA ATOM 2898 CB LYS 362 2.961 30.570 69.070 1.00 22.91 AAAA ATOM 2899 CG LYS 362 2.145 31.200 70.168 1.00 50.86 AAAA ATOM 2900 CD LYS 362 2.145 31.200 70.168 1.00 57.51 AAAA ATOM 2901 CE LYS 362 2.290 32.715 70.157 1.00 57.51 AAAA ATOM 2901 CE LYS 362 1.923 33.278 68.799 1.00 50.87 AAAA ATOM 2902 NZ LYS 362 2.307 34.683 68.711 1.00 22.99 AAAA ATOM 2903 C LYS 362 5.269 30.014 68.202 1.00 16.77 AAAA	ATOM	2893	NH2	ARG	361	8.861	36.704	63.081	1.00 23.47	AAAA
ATOM 2895 O ARG 361 6.926 32.255 68.968 1.00 21.79 AAAA ATOM 2896 N LYS 362 4.681 32.338 68.763 1.00 24.89 AAAA ATOM 2897 CA LYS 362 4.479 30.938 69.125 1.00 28.63 AAAA ATOM 2898 CB LYS 362 2.961 30.570 69.070 1.00 22.91 AAAA ATOM 2899 CG LYS 362 2.145 31.200 70.168 1.00 50.86 AAAA ATOM 2900 CD LYS 362 2.145 31.200 70.168 1.00 57.51 AAAA ATOM 2901 CE LYS 362 2.290 32.715 70.157 1.00 57.51 AAAA ATOM 2901 CE LYS 362 1.923 33.278 68.799 1.00 50.87 AAAA ATOM 2902 NZ LYS 362 2.307 34.683 68.711 1.00 22.99 AAAA ATOM 2903 C LYS 362 5.269 30.014 68.202 1.00 16.77 AAAA	MOTA	2894	С	ARG	361	5.897	32.879	68.714	1.00 26.11	AAAA
ATOM 2896 N LYS 362 4.681 32.338 68.763 1.00 24.89 AAAA ATOM 2897 CA LYS 362 4.479 30.938 69.125 1.00 28.63 AAAA ATOM 2898 CB LYS 362 2.981 30.570 69.070 1.00 22.91 AAAA ATOM 2899 CG LYS 362 2.145 31.200 70.168 1.00 50.86 AAAA ATOM 2900 CD LYS 362 2.290 32.715 70.157 1.00 57.51 AAAA ATOM 2901 CE LYS 362 1.923 33.278 68.799 1.00 50.87 AAAA ATOM 2902 NZ LYS 362 2.307 34.683 68.711 1.00 22.99 AAAA ATOM 2903 C LYS 362 5.269 30.014 68.202 1.00 16.77 AAAA										
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ATOM 2898 CB LYS 362 2.961 30.570 69.070 1.00 22.91 AAAA ATOM 2899 CG LYS 362 2.145 31.200 70.168 1.00 50.86 AAAA ATOM 2900 CD LYS 362 2.290 32.715 70.157 1.00 57.51 AAAA ATOM 2901 CE LYS 362 1.923 33.278 68.799 1.00 50.87 AAAA ATOM 2902 NZ LYS 362 2.307 34.683 68.711 1.00 22.99 AAAA ATOM 2903 C LYS 362 5.269 30.014 68.202 1.00 16.77 AAAA										
ATOM 2899 CG LYS 362 2.145 31.200 70.168 1.00 50.86 AAAA ATOM 2900 CD LYS 362 2.290 32.715 70.157 1.00 57.51 AAAA ATOM 2901 CE LYS 362 1.923 33.278 68.799 1.00 50.87 AAAA ATOM 2902 NZ LYS 362 2.307 34.683 68.711 1.00 22.99 AAAA ATOM 2903 C LYS 362 5.269 30.014 68.202 1.00 16.77 AAAA	MOTA		CA		362					
ATOM 2899 CG LYS 362 2.145 31.200 70.168 1.00 50.86 AAAA ATOM 2900 CD LYS 362 2.290 32.715 70.157 1.00 57.51 AAAA ATOM 2901 CE LYS 362 1.923 33.278 68.799 1.00 50.87 AAAA ATOM 2902 NZ LYS 362 2.307 34.683 68.711 1.00 22.99 AAAA ATOM 2903 C LYS 362 5.269 30.014 68.202 1.00 16.77 AAAA	ATOM	2898	CB	LYS	362	2.981	30.570	69.070	1.00 22.91	AAAA
ATOM 2900 CD LYS 362 2.290 32.715 70.157 1.00 57.51 AAAA ATOM 2901 CE LYS 362 1.923 33.278 68.799 1.00 50.87 AAAA ATOM 2902 NZ LYS 362 2.307 34.683 68.711 1.00 22.99 AAAA ATOM 2903 C LYS 362 5.269 30.014 68.202 1.00 16.77 AAAA						2.145	31.200		1.00 50.86	AAAA
ATOM 2901 CE LYS 362 1.923 33.278 68.799 1.00 50.87 AAAA ATOM 2902 NZ LYS 362 2.307 34.683 68.711 1.00 22.99 AAAA ATOM 2903 C LYS 362 5.269 30.014 68.202 1.00 16.77 AAAA										
ATOM 2902 NZ LYS 362 2.307 34.683 68.711 1.00 22.99 AAAA ATOM 2903 C LYS 362 5.269 30.014 68.202 1.00 16.77 AAAA					•				1 00 50 97	
ATOM 2903 C LYS 362 5.269 30.014 68.202 1.00 16.77 AAAA										
		2902	NZ	LYS					1.00 22.99	
	ATOM	2903	С	LYS	362		30.014	68.202	1.00 16.77	
E135 2 0 1	ATOM	2904	0	LYS	3.62	5.808	29.007	68.647	1.00 22.90	AAAA .

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Figure 16-45

					•				
ATOM	2905	N	GLU	363	5.311	30.355	66.913	1.00 25.24	AAAA
	2906	CA	GLU		6.055	29.577	65.910	1.00 26.29	AAAA
ATOM									
MOTA	2907	CB	GLU		6.207	30.342	64.608	1.00 33.50	AAAA
MOTA	2908	CG	GLU	363	4.999	30.639	63.824	1.00 48.73	AAAA
ATOM	2909	CD	GLU	363	5.368	31.494	62.638	1.00 42.01	AAAA
ATOM -			GLU		6.299	31.087	61.895	1.00 28.50	AAAA
MOTA	· 2911	OE2			4.738	32.558	62.461	1.00 44.91	AAAA
ATOM	2912	С	GLU	363	7.481	29.326	66.349	1.00 19.00	ÄAAA
MOTA	2913	0	GLU	363	8.011	28.218	66.226	1.00 18.66	AAAA
	2914		VAL	364	8.121	30.399	66.790	1.00 20.69	AAAA
ATOM		Ŋ							
MOTA	2915	CA	VAL	364	9.501	30.303	67.219	1.00 23.13	AAAA
ATOM	2916	CB	VAL	.364	10.096	31.681	67.510	1.00 16.98	AAAA
MÓTA	2917	CG1	VAL	364	11.515	31.513	68.010-	1.00 22.32	AAAA
ATOM	2918		VAL	364	10.082	32.548	66.242	1.00 23.99	AAAA
					9.625		68.448	1.00 19.28	
ATOM	2919	C,	VAL	364		29.415			AAAA
ATOM	2920	0	VAL	364	10.507	28.548	68.510·	1.00 20.17	- Aaaa
ATOM	2921	N	LYS	365	8.735	29.600	69.417	1.00 21.11	AAAA
ATOM	2922	CA	LYS	365	8.780	28.768	70.612	1.00 18.15	AAAA
	2923		LYS	365	7.711	29.210	71.626	1.00 25.22	AAAA
ATOM		CB							
MOTA	2924	CG	LYS	365	7.921	30.611	72.167	1.00 32.99	KAAA
MOTA	2925	CD	LYS	365	6.901	30.949	73.253	1.00 36.09	AAAA
MOTA	2926	CE	LYS	365	7.121	32.357	73.790	1.00 28.99	AAAA
	2927	NZ	LYS	365	6.178	32.736	74.882	1.00 38.98	AAAA
ATOM									
MOTA	2928	С	LYS	365	8.574	27.305	70.236	1.00 19.49	AAAA
ATOM	2929	0	LYS	365	9.255	26.417	70.758	1.00 22.04	aaaa
ATOM	2930	N	ASP	366	7.635	27.048	69.327	1.00 22.45	AAAA
ATOM	2931	CA	A.S.P	366	7.386	25.669	68.915	1.00 22.62	AAAA
		CB	ASP	366	. 6.173	25.574	67.967	1.00 21.69	
MOTA	2932								AAAA
MOTA	2933	CG	ASP	366	4.870	25.987	68.634	1.00 27.75	AAAA
MOTA	2934	OD1	ASP	366	4.763	25.890	69.881	1.00 31.01	AAAA
ATOM	2935	OD2	ASP	366	3.938	26.382	67.907	1.00 33.20	AAAA
ATOM	2936	С	ASP	366	8.606	25.034	68.237	1.00 24.53	AAAA
					8.924	23.871	68.480	1.00 21.13	AAAA
MOTA	2937	0	ASP	366					
ATOM	2938	N	THR	367	9.281	25.787	67.380	1.00 26.19	AAAA
ATOM	2939	CA	THR	367	10.462	25.252	66.694	1.00 21.68	AAAA
ATOM	2940	CB	THR	367	11.035	26.301	65.742	1.00 14.56	AAAA.
ATOM	2941	0G1	THR	367	10.085	26.545	64.697	1.00 21.76	AAAA
	2942		THR	367	12.340	25.825	65.138	1.00 19.83	AAAA
ATOM									
ATOM	2943	С	THR	367	11.523	24.822	67.710	1.00 19.02	AAAA
ATOM	. 2944	0	THR	367	12.071	23.717	67.625	1.00 21.79	AAAA
ATOM	2945	71	LEU	368	11.802	25.684	68.683	1.00 18.42	aaaa
ATOM	2946	CA	LEU	368	12.797	25.348	69.700	1.00 21.02	AAAA
	2947	CB	LEU	368	13.148	26.569	70.560	1.00 17.34	AAAA
MOTA								1.00 17.45	
MOTA	2948	CG	LEU	368	14.206	27.518	69.959		AAAA
ATOM	2949		LEU	368	15.525	26.758	69.817	1.00 16.83	AAAA
ATOM	2950	D2	LEU	368	13.756	28.041	68.593	1.00 19.49	AAAA
ATOM	2951	J	LEU	368	12.361	24.189	70.589	1.00 23.17	AAAA
ATOM	2952	Ū	LEU	368	13.203	23.420	71.052	1.00 24.81	AAAA
									AAAA
MOTA	2953	Ŋ	GLU	369	11.059	24.055	70.839	1.00 23.97	
ATOM	2954	CA	GLU	369	10.597	22.929	71.653	1.00 19.36	AAAA
ATOM	2955	CB	GLU	369	9.127	23.113	72.063	1.00 21.81	Aaaa
ATOM	2956	CG	GLU	369	8.913	24.225	73.100	1.00 40.15	AAAA
				369	7.450	24.416	73.487	1.00 49.38	AAAA
MOTA	2957	CD	GLU						
MOTA	2958	OEI	GLU	369	6.806	23.429	73.905	1.00 43.26	AAAA.
ATOM	2959	OE2	GLU	369	6.948	25.558	73.382	1.00 57.31	AAAA
ATOM	2960	С	GLU	369	10.778	21.623	70.859	1.00 24.29	AAAA
	2961	ō	GLU	369	11.172	20.605	71.420	1.00 25.96	AAAA
ATOM							69.560	1.00 22.98	AAAA
ATOM	2962	N_	LYS	370	10.488	21.643			
MOTA	2963	CA	LYS	370	10.665	20.437	68.746	1.00 23.19	AAAA
MOTA	2964	CB .	LYS	370	10.051	20.596	67.347	1.00 26.83	AAAA
ATOM	2965	CG	LYS	370	8.537	20.461	67.287	1.00 36.68	AAAA
	2966	CD	LYS	370	8.056	20.431	65.832	1.00 39.85	AAAA
MOTA								_	AAAA
MOTA	2967	CE	LYS	370	6.567	20.105	65.740	1.00 56.23	
ATCM ·	2968	NZ	LYS	370	6.082	19.996	64.326	1.00 56.10	AAAA
MOTA	2969	С	LYS	370	12.148	20.123	68.602	1.00 31.63	AAAA
•	2970	ō	LYS	370	12.549	18.958	68.587	1.00 36.88	AAAA
MOTA	4210	٠.		3.0		-0.550	•	•	•
•									

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ATOM	297	71 N	ALA	371	12.96	1 21.17	0 68.493	1 1 0	0 26.25	2222
ATOM	297	72 CA	ALA	371	14.40				0 27.33	
MOTA	297	3 CE	ALA	371	15.07			1.0	0 27.33	
ATOM	297		ALA	371	14.98				0 23.70	
ATOM	297		ALA	371					0 26.74	AAAA
					15.89			2 1.0	0 29.52	AAAA
ATOM	297		ALA	372	14.48	4 20.652	2 70.764	1.0	0 24.83	
MOTA	297		ALA	372	14.95	9 20.059	72.012		0 34.24	
ATOM	297	8 CB	ALA	3 72	14.30	5 20.750		-	0 37.17	AAAA
ATOM	297	9 C	ALA	.372	14.66					AAAA
MOTA	298		ALA	372	15.56				0 45.62	AAAA
ATOM	298	-	ALA	373					0 35.52	AAAA
MOTA			•		13.39				0 44:50	AAAA
	298	_		373	13.00		71.892	1.0	0 49.88	AAAA
MOTA	298		ALA	3 7 3	11.50	6 16.681	71.628		0 49.32	AAAA
ATOM	298	4 C	ALA	373	13.80	7 16.072			0 44.64	
ATOM	298	5 0	ALA	373	14.66					AAAA
ATOM	298	סצי	r ALA	373	13.59			-	0 58.19	AAAA
HETATI		-	TAW S	1					0 41.63	AAAA
HETATI		_			36.36				0 13.03	SOLV
			TAW S	2	23.10				0 11.42	SOLV
HETATI			TAW S	3	20.59	4 33.744	61.457	1.0	0 14.73	SOLV
HETATI	4 299	OH2	TAW S	4	31.359	9 16.551			0 19.84	SOLV
HETATN	1 299:	L OH2	TAW S	5	30.389				0 19.94	
HETATN	1 2992	2 OH2	TAW	6	16.92					SOLV
HETATM	4 2993	_	YAW S	7	28.448				0 13.33	SOLV
HETATM			WAT	8					0 14.08	SOLV
HETATM					40.375				0 19.10	SOLV
			WAT	9	18.455			1.00	0 18.81	SOLV
HETATM			WAT	10	26.305			1.00	16.65	SOLV
HETATM			WAT	11	50.145	32.063	58.142		16.53	SOLV
HETATM	1 2998	OH2	WAT	12	45.935	30.996	40.672		25.08	SOLV .
HETATM	1 2999	OH2	WAT	13	26.358		74.179		22.91	
HETATM	3000	OH2	WAT	14	48.727		56.917			SOLV
HETATM			WAT	15	30.244				25.49	SOLV
HETATM		. –	WAT	16			50.165		25.78	SOLV
HETATM					10.615		63.631		22.40	SOLV
		_	WAT	17	18.401		62.704	1.00	21.46	SOLV
HETATM			WAT	18	22.195		60.896	1.00	26.19	SOLV
HETATM			WAT	19	3.278	32.141	65.350	1.00	20.38	SOLV
HETATM		· 0H2	WAT	20	23.643	22.897	59.512		21.27	SOLV
HETATM	3007	OH2	WAT	21	50.287		48.818		19.73	
HETATM	3008	OH2	WAT	22	44.725		46.541			SOLV
HETATM	3009		WAT	23	8.346	30.527			18.74	SOLV
HETÄTM			WAT	24	39.855		49.922		22.33	SOLV
HETATM			WAT			33.795	67.390		20.43	SOLV
				25	7.827	32.763	57.779	1.00	19.24	SOLV
HETATM			WAT	26	45.388	34.567	36.246	1.00	20.86	SOLV
HETATM			WAT	27	47.636	32.244	33.388	1.00	20.41	SOLV
HETATM		OH2	WAT	28	32.514	35.684	41.278	1.00	24.76	SOLV
HETATM	3015	OH2	WAT	29	26.188	15.341	61.913		19.63	SOLV
HETATM	3016	OH2	TAW	30	14.957	43.169	56.333		23.80	
HETATM	3017	OH2		31	24.483	43.556	55.704			SOLV
HETATM	3018	OH2		32	41.141	16.376			27.25	SOLV
HETATM		OH2		33			48.456		25.99	SOLV
HETATM					23.104	17.625	54.086		26.37	SOLV
		OH2		34	51.301	28.602	57.694	1.00	32.78	SOLV
HETATM		OH2		35	51.376	29.469	53.156	1.00	24.27	SOLV
HETATM	3022	OH2	WAT	36	12.518	22.131	49.816		23.60	SOLV
HETATM	3023	OH2	WAT	3 7	6.521	27.442	50.861		25.87	SOLV
METATM	3024	OH2	TAW	38	30.390	33.757	34.190		19.87	
HETATM		OH2		39	8.328	29.586				SOLV
HETATM		OH2		40			62.062		32.01	SOLV
					30.180	24.235	30.724		22.61	SOLV
HETATM	•	OH2		41	44.521	30.663	38.395	1.00	27.52	SOLV
HETATM		OH2 (42	30.981	18.043	41.186	1.00	23.45	SOLV
HETATM		он2 (TAK	43	14.632	37.127	73.830		29.36	SOLV
HETATM :	3030	OH2 ;	JAT	44	39.332	25.953	72.230		21.87	SOLV
HETATM :	3031	OH2 V		45	7.597	37.592	51.896		39.62	SOLV
HETATM		OH2 V		46	15.027	18.079				_
HETATM		OH2 V		47	11.076		54.827		26.65	SOLV
						45.493	66.435		38.18	SOLV
HETATM		OH2 V		48	42.124	18.055	37.233		28.62	SOLV
ETATM		OH2 W		49	48.736	25.764	64.149	1.00	31.88	SOLV
ETATM	3036	OH2 W	<i>I</i> AT	5 0	50.383	27.254	54.972	1.00	24.36	SOLV
							-			

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HETATN	4 3037	CH	2 WAT	51	48.659	36.025	68.226	1.00 33.89	SOLV
HETATN	4 3038	OH	2 WAT		36.998	27.228	71.440	1.00 21.03	
									SOLV
HETATI			2 WAT		41.303	16.309	55.307	1.00 32.23	SOLV
HETATI	13040	OH	2 WAT	54	33.242	39.524	49.454	1.00 29.77	SOLV
HETATM	1 3041	OH	2 WAT	55	45.004	25.973	35.031	1.00 21.59	SOLV
HETATM			2 WAT	56	19.039	25.829		1.00 33.48	
									SOLV
HETATM	1 3043	OH	2 WAT	5 7	17.922	35.542	50.154	1.00 37.51	SOLV
HETATM	1 3044	OH	2 WAT	58	10.409	26.864	73.166	1.00 26.54	SOLV
HETATM	13045	OH	2 WAT	59	11.835	22,805	59.408	1.00 20.83	SOLV
			-						
HETATM			2 WAT	60.	18.254	48.699	53.224	1.00 28.41	. SOLV
HETATM	3047	OH	2 WAT	61	10.426	26.647	60.447	1.00 32.72	SOLV
HETATM	1 3048	OH	2 WAT	62	21.304	55.086	63.510	1.00 28.84	SOLV
HETATM		. Un.	2 WAT	63	32.532	51.211	45.469	1.00 32.48	
									SOLV
HETATM			2 WAT	64	22.658	61.079	57.420	1.00 27.32	SOLV
HETATM	1 3051	OH:	2 WAT	65	16.734	24.334	74.721	1.00 27.44	SOLV
HETATM	3052	OH:	2 WAT	66	32.758	37.824	54.391	1.00 25.07	SOLV
HETATM			TAW S	67	11.142	25.859	49.706	1.00 29.66	
									solv
HETATM			TAW S	68	24.192	15.261	53.236	1.00 30.21	SÓLV
HETATM	3055	OH	TAW S	69	19.816	17.916	66.357	1.00 30.50	SOLV
HETATM	3056	OH	TAW S	70	50.347	23.975	53.197	1.00 28.08	SOLV
HETATM			TAW S	71	50.258	30.918	51.113		
								1.00 20.19	SOLV
HETATM	3058		TAW S	72	21.047	17.624	68.693	1.00 41.23	SOLV
HETATM	3059	OH	TAW	73	26.782	33.756	49.995	1.00 25.80	SOLV
HETATM	3060	OH2	TAW S	74	12.570	43.844	64.441	1.00 31.03	SOLV
HETATM			WAT	75	35.555	41.287	50.852	1.00 24.03	
									SOLV
HETATM	3062		NAT	76	27.764	18.231	61.827	1.00 18.28	SOLV
HETATM	3063	OH2	YAT	7 7	26.715	29.236	38.391	1.00 23.18	SOLV
HETATM	3064	OH2	WAT	78	21.461	23.245	48.872	1.00 23.80	SOLV
HETATM			WAT	79	49.246	28.263	65.477	1.00 21.52	
									SOLV
HETATM	_		WAT	80	31.785	13.301	69.606	1.00 31.11	SOLV
HETATM	3067	OH2	WAT	81	49.811	34.740	59.229	1.00 31.76	SOLV
HETATM	3068	OH2	WAT	82	45.670	33.188	42.470	1.00 23.13	SOLV
HETATM			WAT	83	9.408	39.751	55.872	1.00 31.53	SOLV
HETATM			WAT	84 -	35.166	35.878	29.899	1.00 37.32	SOLV
HETATM	3071	OH2	WAT	85	41.927	22.970	73.694	1.00 44.07	SOLV
HETATM	3072	OH2	WAT	86	22.125	34.577	49.199	1.00 44.65	SOLV
HETATM	3073	OH2	WAT	87	43.984	33.541	37.965	1.00 24.88	SOLV
HETATM			WAT	88	11.997	17.962	56.312	1.00 34.85	
									SOLV
HETATM	3075		WAT	89	42.194	14.737	59.766	1.00 25.91	SOLV
HETATM	3076	OH2	WAT	90	49.313	24.200	41.684	1.00 29.29	SOLV
HETATM	3077	OH2	WAT	91	48.504	33.595	61.519	1.00 30.32	SOLV
HETATM			WAT	92	24.773	18.356	33.365	1.00 53.13	SOLV
					35.160				
HETATM			WAT	93		35.656	47.470	1.00 41.41	SOLV
HETATM	3080	OH2	WAT	94	44.682	36.658	39.962	1.00 29.24	SOLV
HETATM	3081	OH2	WAT	95	. 9.576	41.033	52.549	1.00 51.83	· SOLV
HETATM	3082	OH2	WAT	<u>~6</u>	47.199	20.112	42.102	1.00 40.39	SOLV
HETATM	-		WAT	7ر ٠	49.254	26.331	59.641	1.00 37.03	
									SOLV
HETATM			TAW	_ 8	26.808	37.600	38.172	1.00 28.74	SOLV
HETATM	3085	OH2	WAT	99	40.749	14.572	64.635	1.00.33.42	SOLV
HETATM	3086	OH2	WAT	100	24.850	44.161	47.775	1.00 27.89	SOLV
			WAT	101	34.326			1.00 42.22	
HETATM									SOLV
HETATM	3088		WAT	102	30.226	34.544	52.026	1.00 30.77	SOLV
HETATM	3089	OH2	WAT.	103	47.824	39.054	78.097	1.00 52.16	SOLV
HETATM			WAT	104	19.665	18.953	47.438	1.00 51.70	SOLV
					46.857		46.232		
HETATM			WAT	105		36.525		1.00 23.65	SOLV
HETATM		OH2	WAT	106	48.069	19.460	67.360	1.00 37.56	SOLV
HETATM	3093	OH2	WAT	107	15.553	56.850	61.838	1.00 46.95	SOLV
HETATM		OH2		108	44.026	19.119	70.671	1.00 39.55	SOLV
					8.139	42.064		1.00 42.61	
HETATM			WAT	109					SOLV
HETATM	3096	OH2	WAT	110	50.624	36.591	65.779	1.00 31.59	SOLV
HETATM		OH2	WAT	111	51.398	26.073	61.043	1.00 49.09	SOLV
HETATM		OH2		112	26.174	33.692	33.551	1.00 36.61	SOLV
					23.545			1.00 24.34	
HETATM		OH2		113		20.203	53.001		SOLV
HETATM	3100	OH2		114	9.083	42.965	57.697	1.00 33.65	SOLV
HETATM	3101	OH2	TAW	115	8.442	39.898	64.594	1.00 31.21	SOLV
HETATM		OH2		116 .	15.219	35.897	51.951	1.00 26.59	SOLV
STATE	- 1 UE	~				JJ. JJ.			

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HETAT	M 310	3 OI	12 WAT	117	1	15.417	38.43	8 50.473	1.0	0 34.46	SOLV
HETAT	M 310	4 01	12 WAT	118		40.757				0 29.12	
HETAT	M 310		12 WAT			27.717				0 23.12	SOLV
HETAT			12 WAT			8.612				0 28.17	SOLV
HETAT			I2 WAT			13.198			1 1 0	0 38.56	
HETAT			2 WAT						1.0	0 26.31	SOLV
						4.188			1.0	0 29.81	SOLV
HETAT			2 WAT			0.736			1.0	0 32.40	SOLV
HETAT			2 WAT			1.302			1.0	0 30.84	SOLV
HETAT	M 311:	r oh	2 WAT	125	3	6.895	21.26	4 34.198	3 1.0	0 34.67	SOLV
HETATI	M 3112	OH S	2 WAT	126	4	7.474	22.25	2 67.427	1.0	0 34.35	SOLV
HETATI	M 3113	OH	2 WAT	127		7.178	25.93	6 64.063	1.0	0 31.77	SOLV
HETATI	4 3114	OH	2 WAT	128	3	6.362	66.64		1.0	0 36.88	SOLV
HETATI	x 3115	ОН	2 WAT	129		2.486				0. 26.61	
HETATI	4 3116		2 WAT	130		8.432				0. 20.01	SOLV
HETATI			2 WAT	131		7.644				0 37.33	SOLV
HETATI			2 WAT			0.273					SOLV
HETATI			2 WAT	133		7.518				0 37.33	SOLV
HETATI			2 WAT	134			46.197			0 45.42	SOLV
										0 28.02	SOLV
HETATI			2 WAT	135		1.501				0 32.78	SOLV
HETAT			2 WAT	136		5.898				0 43.47	SOLV
HETATI			Z WAT	137		6.300				0 30.37	SOLV
HETATM			TAW S	138		1.148	36.946			0 46.34	SOLV
HETATI			2 WAT	139		1.525	53.761		1.0	38.27	SOLV
HETATM			TAW S	140	2:	1.603	54.580	68.690		33.10	SOLV
HETAT			TAW S	141	10	0.191	29.237	60.325	1.00	30.24	SOLV
HETATM		OH2	TAW S	142	10	6.951	18.120	66.901		40.85	SOLV
HETATK	1 3129	OH2	TAW S	143	. 4	4.943	24.912			49.13	SOLV
HETATM	3130	OH2	TAW S	144	10	0.711	25.291	58.177		30.72	SOLV
HETATM	3131	OH2	YAW S	145	3 (0.815	43.398	36.040		42.23	SOLV
HETATM	3132	OH2	YAW S	146		1.763	24.512			28.31	SOLV
HETATM	3133	OH2	WAT -			1.788	33.122			26.15	SOLV
HETATM			WAT	148		1.531	44.741			27.99	SOLV
HETATM			WAT	149		0.938		60.422			
HETATM			WAT	150		1.860	47.932			38.20	SOLV
·HETATM			WAT	151		7.336	37.304			18.89	SOLV
HETATM			WAT	152		3.680				33.58	SOLV
HETATM			WAT	153		1.441	35.535			26.89	SOLV
HETATM			WAT				16.097			48.33	SOLV
				154		343	18.124			36.28	SOLV
HETATM			WAT	155		7.765	37.948			48.41	SOLV
HETATM			WAT	156		.329	31.169			25.33	SOLV
HETATM			TAW	157		.028	24.554			41.54	SOLV
HETATM	_		WAT	158		.888	15.082		1.00	28.76	SOLV
HETATM			TAW	159		.886	20.780		1.00	51.03	SOLV
HETATM			WAT	160		.962	49.969		1.00	35.04	SOLV
HETATM			WAT	161	14	.696	15.261		1.00	55.47	SOLV
HETATM			WAT	162	14	.915	18.181	64.866	1.00	42.00	SOLV
HETATM	3149		WAT	163		.608	49.029	52.612	1.00	47.32	. SOLV
HETATM	3150	OH2	WAT	164	52	.566	30.906	57.612	1.00	36.71	SOLV
HETATM	3151	OH2	WAT	165	23	.699	27.331	77.729	1.00	32.22	SOLV
HETATM	3152	OH2	TAW	166		.971	59.046	63.272	1.00	43.05	SOLV
HETATM	3153	OH2	WAT	167	46	.053	45.927	52.876	1.00	33.66	SOLV
HETATM	3154	OH2	WAT	168	42	.780	49.151	58.106		44.63	SOLV
HETATM	3155	OH2	WAT	169		.100		72.183		45.43	SOLV
HETATM		OH2		170		.677	60.998			34.51	SOLV
HETATM		OH2		171		.336	45.674	45.578		55.85	
HETATM		OH2		172		.481	18.266				SOLV
HETATM		OH2				.112	18.147	49.018		32.73	SOLV
HETATM		OH2		173 174			43 340	31.404		49.94	SOLV
				174 .		.874	43.142	70.985		32.89	SOLV
HETATM		OH2		175		.517	17.884	33.278		42.20	SOLV
HETATM		OH2		176		.330	54.886	50.466		40.74	SOLV
HETATM		OH2		.177		400	51.087	74.689		38.56	SOLV
HETATM		OH2		178		971	27.079	67.130	1.00	44.49	SOLV
HETATM		OH2		179	7.	.933	23.412	54.691	1.00	42.84	SOLV
HETATM	3166	OH2	TAW	180	33.	498	47.596	73.612	1.00	35.99	SOLV
HETATM	3167	OH2	TAW	181	26.	016	19.583	44.954		51.31	SOLV
HETATM .	3168	OH2	WAT	182	40.	139	17.026	74.920		43.64	SOLV
				• •			•			-	

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HETATM 3169	OH2 WAT	183	10.441	42.659	62.744	1.00 34.51	SOLV
HETATM 3170	OH2 WAT	184	2.095	34.482	65.810	1.00 36.49	SOLV
HETATM 3171	OH2 WAT	185	45.749	18.286	51.615	1.00 28.19	SOLV
HETATM 3172	OH2 WAT	186	25.771	38.332	76.707	1.00 45.53	SOLV
HETATM 3173	OH2 WAT	187	7.228	40.382	57.542	1.00 48.91	SOLV
HETATM 3174	OH2 WAT	188	42.972	52.824	67.739	1.00 39.99	SOLV
HETATM 3175	OH2 WAT	189	20.137	13.189	73.277	1.00 44.91	SOLV
HETATM 3176	OH2 WAT	190	48.945	19.193	47.581	1.00 52.88	SOLV
HETATM 3177	OH2 WAT	191	14.549	34.547	47.665	1.00 49.15	SOLV
HETATM 3178	OH2 WAT	192	31.765	20.567	26.536	1.00 42.23	SOLV
HETATM 3179	OH2 WAT	193	9.784	39.303	74.222	1.00 32.10	SOLA
HETATM 3180	OH2 WAT	194	28.865	12.481	52.375	1.00 50.98	SOLV
HETATM 3181	OH2 WAT	195	- 24.030	12.804	70.409-	1.00 52.43	SOLV
HETATM 3182	OH2 WAT	196	47.209	39.53€	50.698	1.00 43.03	SOLV
HETATM 3183	OH2 WAT	197	35.618	18.114	27.306	1.00 41.11	SOLV
HETATM 3184	OH2 WAT	198	23.625	48.145	43.853	1.00 48.20	~ zóra
HETATM 3185	OH2 WAT	199	37.090	59.044	54.185	1.00 34.99	SOLV
HETATM 3186	OH2 WAT	200	34.478	12.208	59.080	1.00 36.58	SOLV
HETATM 3187	OH2 WAT	201	22.142	29.583	76.228	1.00 33.95	· SOLV
HETATM 3188	OH2 WAT	202	13.608	42.619	53.973	1.00 40.44	SOLV
HETATM 3189	OH2 WAT	203	42.647	18.701	72.526	1.00 55.64	SOLV
HETATM 3190	OH2 WAT	204	37.005	35.993	77.480	1.00 34.82	SOLV
HETATM 3191	OH2 WAT	205	34.154	20.512	33.327	1.00 31.00	SOLV
HETATM 3192	OH2 WAT	206	37.264	57.546	47.642	1.00 49.58	SOLV
HETATM 3193	OH2 WAT	207	17.924	35.195	79.003	1.00 38.45	SOLV
HETATM 3194	OH2 WAT	208	51.172	31.581	62.378	1.00 35.37	SOLV
HETATM 3195	OH2 WAT	209	50.503	36.726	79.224	1.00 39.95	SOLV
HETATM 3196	OH2 WAT	210	18.382	13.162	63.852	1.00 52.08	SOLV
HETATM 3197	OH2 WAT	211	27.245	8.351	55.199	1.00 39.12	SOLV
HETATM 3198	OH2 WAT	212	18.354	13.545	59.540	1.00 30.15 1.00 36.69	SOLV
HETATM 3199	OH2 WAT	213	49.088	51.744	63.388 50.871	1.00 38.89	SOLV
HETATM 3200	OH2 WAT	214	23.251	33.160	50.871	1.00 42.11	SOLV
HETATM 3201	OH2 WAT	215	12.989	35.073	43.239	1.00 37.93	SOLV
HETATM 3202	OH2 WAT	216	24.414 24.690	44.460 47.590	73.117	1.00 34.17	SOLV
HETATM 3203	OH2 WAT	217	19.844	17.949	81.360	1.00 40.74	SOLV
HETATM 3204	OH2 WAT	218	40.169	27.215	74.247	1.00 37.83	SOLV
HETATM 3205	OH2 WAT	219	38.737	39.516	73.171	1.00 49.20	SOLV
HETATM 3206	CH2 WAT	220 221	50.628	21.408	46.879	1.00 45.57	SOLV
HETATM 3207	OH2 WAT	221	35.436	43.288	75.660	1.00 37.33	SOLV
HETATM 3208	OH2 WAT	222	34.390	16.963	55.285	1.00 35.10	SOLV
HETATM 3209	OH2 WAT	223	21.800	35.454	34.475	1.00 46.29	SOLV
HETATM 3210	OH2 WAT	225	15.751	40.989	46.787	1.00 62.75	SOLV
HETATM 3211	OH2 WAT	225	23.844	48.662	66.295	1.00 38.35	SOLV
HETATM 3212	OH2 WAT	227	47.225	20.562	55.117	1.00 49.99	SOLV
HETATM 3213	OH2 WAT	228	23.426	19.272	50.565	1.00 30.07	SOLV
HETATM 3214	UNA MAI	440	20.720	,_,	24.434		_

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					T.	ignic 17-	-1					
					Residue	∦ X	Y	Z	occ.	В	Segment	: ID
MOTA	1		ALA	A	2	43.739	36.862	75.052	1.00	64.01	6	
MOTA	2		AĿA	A	2	44.405	38.106	72.971	1.00	60.02	6	
MOTA	3		ALA		2	43.251	38.536	72.908	1.00	57.94	8	
ATOM	4		ALA		2	46.142	37.179	74.497		62.88	7	•
MOTA	5		АĻА		2	44.776	36.966	73.923	1.00	63.02	6	
MOTA	6		LYS	A	3	45.398	38.588	72.233	1.00	55.40	7	
ATOM	7		LYS		3	45.196	39.671	71.287	1.00	53.02	6	
ATOM	8		LYS		3	46.443	39.830	70.421	1.00	53.11	6	
ATOM	9		LYS			47.703	40.093	71.217	1.00	57.36	6	
MOTA	10		LYS		3	48.941	39.976	70.349		60.94	6	
ATOM .	11		LYS		3	48.909	40.957	69.196	1.00	63.48	6	
ATOM	12		LYS		3	50.075	40.765	68.294		66.87	7	
MOTA	13	C	LYS		3	43.986	39.401	70.399		50.49	6	
MOTA	14	0	LYS		3	43.691	38.255	70.063		52.50	8	
MOTA	15		VAL		4	43.281	40.464	70.034		45.96	7	
MOTA	16	CA	VAL		4	42.122	40.352	69.167		41.16	6	
ATOM	17	CB	VAL		4	40.983	41.272	69.638		41.53	6	
ATOM	18		VAL		4	39.734	41.028	68.797		40.07	6	
ATOM	19		VAL		4	40.705	41.033	71.115		38.31	6	
ATOM	20	С	VAL		4	42.619	40.796	67.796		39.96	6	
ATOM.	21	0	VAL		4	43.123	41.914	67.645		39.15	8	
ATOM	22	N	LYS		5 5	42.486	39.916	66.807		36.24	7	
MOTA	23 24	CA	LYS		5	42.956	40.186	65.449		35.66	6 .	
ATOM	25	CB CG	LYS		5	43.930 45.197	39.088 38. <i>9</i> 78	65.024 65.860		37.33	6	
ATOM ATOM	26	CD	LYS		5	46.113	40.179	65.659		38.24	6	
ATOM	27	CE	LYS		5	47.436	39.957	66.369		37.46	6 6	
ATOM	28	NZ	LYS		5	48.345	41.121	66.245		35.63	7	
ATOM	29	C	LYS		5	41.840	40.254	64.415		34.40	6	
ATOM	30	ō	LYS		5	40.788	39.641	64.588		33.92	8	
MOTA	31	N	LEU		6	42.082	40.983	63.329		32.52	7	
ATOM	32	CA	LEU		6	41.097	41.094	62.253		33.64	6	
MOTA	33	CB	LEU	A	6	40.589	42.532	62.114		31.83	6	
MOTA	34	CG	LEU	Α	6	39.346	42.823	61.248		32.93	6	
MOTA	35	CD1	LEU	Α	6	39.356	44.295	60.899	1.00	28.95	6	
ATOM	36	CD2	LEU	Α	6	39.336	42.031	59.964	1.00	32.87	· 6	
MOTA	37	С	LEU	A	6	41.802	40.721	60.955	1.00	35.09	6	
MOTA	38	0	LEU	A	6	42.631	41.491	60.468	1.00	36.93	8	
MOTA	39	N	ILE		7 .	41.494	39.561	60.382	1.00	35.52	7	
ATOM	40	CA	ILE		7	42.145	39.199	59.129		35.14	6	
ATOM	41	CB	ILE		7	42.062	37.711	58.850	1.00		6	
ATOM	42		ILE		7	42.731	37.409	57.517		32.87	5 .	
ATOM	43		ILE		7	42.746	36.941	59.975		33.32	6	
ATOM	44		ILE		7	42.744	35.451	59.755	1.00		6	
ATOM	45	C	ILE		7	41.487	39.935	57.971	1.00		6	
ATOM	46 47	N O	GLY .		7 8	40.258 42.304	39.933 40.563	57.855 57.124	1.00		8	
ATOM ATOM "	48	CA	GLY .		8	41.771	41.305	55.994	1.00		. 7	
ATOM	49	C	GLY .		8	42.809	41.939	55.079	1.00		6 6	
ATOM	50	0	GLY .		8		41.827	55.321	1.00		8	
ATOM	51	N	THR		9	42.335	42.622	54.033	1.00		7	
ATOM	52	CA	THR		9	43.212	43.268	53.057	1.00		6.	
ATOM	53	CB	THR		9	44.132	42.210	52.390	1.00		6	
ATOM	54		THR 2		9	44.754	42.771	51.230	1.00		8	
ATOM	55		THR A		وَ	43.332	40.972	52.001	1.00		6	
ATOM	56	С	THR A	À	9	42.447	44.045	51.970	1.00		6	
ATOM	57		THR A		9	41.434	43.569	51.452	1.00		8	
ATOM	58		LEU 2		10	42.939	45.238	51.628	1.00		7	
ATOM	59		LEU ?		10	42.304	46.077	50.609	1.00		6	
ATOM	60		LEU 2		10	43.026	47.418	50.456	1.00		6	
ATOM	61		LEU A		10	42.836	48.506	51.510	1.00	39.68	6	
ATOM	62		LEU A		10	41.343	48.830	51.594	1.00		6	
ATOM	63	CD2	LEU A		10	43.382	48.057	52.857	1.00		6	
ATOM	é÷	С	LEU A	1	10	42.238	45.432	49.239	1.00		б	
ATOM	65		LEU A		10	41.462	45.863	48.381	1.00	42.08	8	
MOTA	66	N .	ASP A		11	43.052	44.408	49.025	1.00	43.51	7	

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Figure 17-2

ATOM	67	CA	ASP		11		43.071	43.731	47.737	1.00 47.27	6
ATOM	68		ASP		11		44.250		47.694	1.00 51.03	6
MOTA	69	CG	ASP		11		45.579	43.479	47.858	1.00 54.10	6
MOTA	70		1 ASP		11		45.944	44.282	46.975	1.00 \$5.93	8
ATOM	71	OD:	2 ASP	A	11		46.255	43.251	48.879	1.00 57.79	8
ATOM	72	С	ASP	A	11		41.756	43.016	47.423	1.00 46.36	6
ATOM	73.	0	ASP		11		41.472	42.702	46.266	1.00 43.49	8
	74	N	TYR		12		40.954	42.767	48.456		7
ATOM										1.00 46.80	
MOTA	75	CA	TYR		12		39.654	42.116	48.284	1.00 45.92	6
ATOM	76	CB	TYR	. A	12		38.953	41.942	49.638	1.00 41.38	6
MOTA	77	CG	TYR	A	12	-	39.358	40.697	50.390	1.00 38.82	6
ATOM	78	CD:	1 TYR	a	12		39.531	40.720	51.775	1.00 37.95	6
ATOM	79	CE:			12		39.869	39.560	52.476	1.00 36.18	6
	80	CD			12		39.533	39.479	49.721	1.00 37.69	
MOTA											6
ATOM	81	CE			12		39.868	38.316	50.415	1.00 35.83	6
MOTA	82	CZ	TYR		12		40.032	38.365	51.787		6
MOTA	83	ОН	TYR	A	12		40.339	37.216	52.470	1.00 36.31	8
ATOM	84	С	TYR	À	12		38.786	42.966	47.378	1.00 46.56	6
ATOM	85	0	TYR		12		37.821	42.476	46.791	1.00 47.38	8
ATOM	86	N	GLY		13		39.138	44.247	47.278	1.00 47.28	7
								45.164			
ATOM	87	CA	GLY		13		38.385		46.442	1.00 46.53	6
ATOM	88	С	GLY		13		38.650	44.934	44.968	1.00 45.60	6
ATOM	89	0	GLY	A	13		37.895	45.401	44.117	1.00 43.68	8
MOTA	90	N	LYS	A	14		39.725	44.210	44.672	1.00 46.52	7
ATOM	91	CA	LYS		14		40.112	43.908	43.296	1.00 47.28	6
ATOM	92	CB	LYS		14		41.629	43.748	43.201	1.00 50.22	6
ATOM	93	CG	LYS		14		42.396	45.044	43.307	1.00 57.12	6
									42.161		
MOTA	94	CD	LYS		14		42.038	46.004		1.00 63.60	6
ATOM	95	CE	LYS		14		42.349	45.422	40.768	1.00 66.65	6
- MOTA	96	NZ	LYS		14		41.529	44.220	40.387	1.00 67.70	7
MOTA	97	С	LYS	A	14		39.460	42.643	42.769	1.00 44.18	6
ATOM	98	0	LYS	A	14		39.564	42.325	41.585	1.00 40.33	8
ATOM	99	N	TYR	A	15		38.790	41.926	43.661	1.00 43.25	7
ATOM	100	CA	TYR		15		38.145	40.665	43.317	1.00 43.18	6
	101	CB	TYR		15	•	38.789	39.547	44.142	1.00 36.88	6
MOTA											
ATOM	102	CG	TYR		15		40.302		44.053	1.00 32.96	6
MOTA	103	CD1			15		41.084	39.107	45.108	1.00 30.90	6
ATOM	104	CE1	TYR	Α	15		42.476	39.144	45.035	1.00 30.94	6
ATOM	105	CD2	TYR	A	15		40.952	40.049	42.912	1.00 33.01	6
MOTA	106	CE2	TYR	A	15		42.341	40.092	42.826	1.00 29.68	6
ATOM	107	CZ	TYR		15		43.098	39.639	43.890	1.00 30.99	6
ATOM	108	ОН	TYR		15		44.471	39.673	43.809	1.00 28.02	8
	109	c	TYR		15		36.661	40.778	43.621	1.00 45.56	6
ATOM											
ATOM	110	0	TYR		15		36.149	40.153	44.552	1.00 45.22	8
ATOM	111	N	ARG		16		35.981	41.599	42.830	1.00 48.81	7
MOTA	112	CA	ARG	A	16		34.553	41.819	42.999	1.90 53.22	6
ATOM	113	CB	ARG	A	16		34.193	43.263	42.654	1 [.] J0 57.11	5
ATOM	114	CG	ARG	A	16		34.852	44.330	43.490	·1 70 61.66	6
ATOM	115	CD	ARG		16		34.280	44.408	44.886	1.00 67.04	6
ATOM	116	NE	ARG		16		34.798	45.590	45.569	1.00 73.59	7
	117	CZ	ARG		16			46.837	45.141	1.00 75.03	6
ATOM .							34.612				
	118				16		33.917	47.065	44.033	1.00 73.03	7
ATOM	119	NH2	ARG	A	16		35.142	47.856	45.808	1.00 75.79	7
MOTA	120	С	ARG	A	16		33.757	40.903	42.080	1.00 51.79	6
MOTA	121	0	ARG	A	16		34.192	40.593	40.970	1.00 50.89	8
ATOM	122	N	TYR		17		32.596	40.463	42.552	1.00 50.19	
	123	CA	TYR		17		31.737	39.634	41.733	1.00 49.60	6
ATOM										1.00 45.80	6
ATCM	124	CB	TYR		17		30.534	39.119	42.528		
MOTA	125	CG	TYR		17		30.803	37.894	43.365	1.00 42.01	6
ATOM	126	CD1	TYR	A	17		31.689	37.932	44.438	1.00 42.45	6
ATOM	127	CEI	TYR	A	17		31.960	36.780	45.193	1.00 43.36	6
ATOM	128	CD2	TYR		17		30.185	36.680	43.062	1.00 39.89	6
	129	CE2	TYR		17		30.443	35.526	43.803	1.00 40.56	6
ATOM							31.333		44.869	1.00 41.98	6
ATOM	130	CZ	TYR .		17			35.578			
ATOM	131	OH	TYR .		17		31.600	34.438	45.598	1.00 34.66	8
ATOM	132	С	TYR .	À,	17		31.245	40.547	40.622	1.00 51.88	6

70/263. Figure 17-3

ATOM	133	0	TYR A	17		31.332	41.772	40.726	1.00 47.86	8
ATOM	134	N	PRO A			30.730	39.964	39.534	1.00 54.38	7
			PRO A			30.548	38.545		1.00 54.21	
ATOM	135	CD						39.190		6
ATOM	136	CA.				30.243	40.809	38.449	1.00 56.43	6
ATOM	137	CB	PRO A			29.601	39.792	37.496	1.00 56.84	6
ATOM -	138	CG	PRO A	. 18		29.260	38.613	38.426°	1.00 56.46	6
MOTA	139	С	PRO A	. 18		29.273	41.891	38.932	1.00 58.74	6
MOTA	140	0	PRO A	. 18		28.791	41.861	40.066	1.00 55.72	8
ATOM	141	N	LYS A			29.017	42.851	38.052	1.00 62.10	7
ATOM	142	CA	LYS A			28.127	43.973	38.314	1.00 64.85	6
						27.972	44.781			
ATOM	143	CB	LYS A					37.022	1.00 69.74	6
MOTA	144	CG	LYS A			28.008	43.925	35.740	1.00 74.99	5
MOTA	145	CD	LYS A		-	26.895	42.881	35.668	- 1.00 78.18	6
MOTA	146	CE	LYS A	19		26.981	42.010	34.420	1.00 80.24	· 6
MOTA	147	NZ	LYS A	19		25.867	41.010	34.361	1.00 81.13	7
ATOM	148	С	LYS. A	19		26.750	43.619	38.869	1.00 64.77	6
ATOM	149	0	LYS A	19		26.414	43.961	40.001	1.00 66.50	.8
MOTA	150	N	ASN A	20		25.957	42.933	38.062	1.00 63.75	ž
	151	CA	ASN A	20		24.612	42.556	38.439	1.00 62.96	6
MOTA					-					
ATOM	152	CB	ASN A	20		23.870	42.031	37.208	1.00 67.42	6
ATOM	153	CG	ASN A	20		22.392	41.833	37.459	1.00 72.29	
ATOM	154		ASN A	20		21.666	42.785	37.772	1.00 75.25	. 8
ATOM	155	ND2	ASN A	20		21.931	40.594	37.322	1.00 74.38	7
ATOM	156	С	ASN A	20		24.602	41.512	39.547	1.00 61.30	6
ATOM	157	0	ASN A	20		23.629	40.773	39.698	1.00 61.49	8
ATOM	158	N	HIS A	21		25.681	41.444	40.321	1.00 57.30	7
ATOM	159	CA	HIS A	21		25.755	40.480	41.418	1.00 54.68	6
ATOM	160	CB	HIS A	21		27.071	39.700	41.373	1.00 52.63	6
ATOM	161	CG	HIS A	21		27.058	38.449	42.195	1.00 49.39	6
	162		HIS A	21		27.336	38.236	43.503		6
ATOM									1.00 49.39	
ATOM	163		HIS A	21		26.664	37.229	41.686	1.00 48.27	7
ATOM	164		HIS A	21		26.704	36.320	42.643	1.00 48.16	6
ATOM	165		HIS A	21		27.108	36.905	43.757	1.00 47.33	7
MOTA	166	С	HIS A	21		25.664	41.215	42.760	1.00 52.89	6
MOTA	167	0	HIS A	21		26.295	42.256	42.947	1.00 52.52	8
ATOM	168	N	PRO A	22		24.880	40.679	43.713	1.00 50.81	7
MOTA	169	CD	PRO A	22		24.076	39.444	43.661	1.00 48.50	6
MOTA	170	CA	PRO A	22		24.734	41.310	45.029	1.00 48.02	6
ATOM	171	CB	PRO A	22		23.860	40.308	45.783	1.00 47.45	6
ATOM	172	CG	PRO A	.22		22.990	39.754	44.667	1.00 47.76	6
ATOM	173	C	PRO A	22		26.074	41.558	45.727	1.00 46.48	6
	174	ō	PRO A	22		26.164	42.405	46.615	1.00 45.69	8
MOTA										7
ATOM	175	N	LEU A	23		27.107	40.816	45.318	1.00 44.97	
ATOM	176	CA	LEU A	23		28.441	40.949	45.906	1.00 41.31	6
ATOM	177	CB	LEU A	23		29.076	39.569	46.131	1.00 39.22	6
MOTA	178	CG	LEU A	23		28.264	38.561	46.953	1.00 37.71	6
ATOM	179	CD^{r}	LEŲ A	23		29.075	37.288	47.157	1.00 35.07	6
ATOM	180	CDI	LEU A	23		27.896	39.165	48.292	1.00 36.90	6
ATOM	181	С.	LEU A	23		29.334	41.789	45.003	1.00 40.14	6
ATOM	182	0	LEU A	23		30.556	41.614	44.951	1.00 39.00	8
ATOM	183	N	LYS A	24		28.706	42.705	44.284	1.00 39.67	7
ATOM	184	CA	LYS A	24		29.430	43.590		.1.00 42.88	6
	185	CB	LYS A	24		28.480	44.120	42.323	1.00 40.24	6
ATOM									1.00 44.08	6
ATOM	186	CG	LYS A	24		28.949	45.362	41.610		
ATOM	187	CD	LYS A	24		28.247	46.600	42.166	1.00 44.47	6
ATCM	188	CE	LYS A	24		26.732	46.492	41.968	1.00 43.23	6
ATOM	189	NZ	LYS A	24		25.989	47.717	42.362	1.00 39.79	7
ATOM	190	С	LYS A	24		30.031	44.723	44.217	1.00 43.70	6
ATOM	191	0	LYS A	24		31.027	45.332	43.817	1.00 47.22	8
ATOM	192	N	ILE A	25		29.431	44.976	45.378	1.00 42.27	7
ATCM	193	CA	ILE A	25		29.870	46.035	46.289	1.00 39.86	6
	194	CB	ILE A	25		28.763	46.407	47.306	1.00 37.72	6
ATOM				25		27.539	46.953	46.580	1.00 39.67	6
ATOM	195		ILE A			28.410			1.00 35.25	6
ATCM	196		ILE A	25			45.168	48.145		6
ATOM	197		ILE A	25		27.301	45.368	49.151	1.00 31.74	
ATCM	198	С	ILE A	25		31.078	45.625	47.112	1.00 40.23	6

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ATOM	199	0	ILE A	25		31.419	44.441	47.198	1.00 38.90	8
	200	N	PRO A			31.762	46.616	47.709		7
MOTA									1.00 40.18	
ATOM	201	CD	PRO A	_		31.523	48.051	47.533	1.00 40.58	6
ATOM	202	CA	PRO A			32.939	46.437	48.558	1.00 38.31	6
MOTA	203	CB	PRO A	. 26		33.478	47.860	48.688	1.00 37.14	6
MOTA	204	CG	PRO A	26		32.940	48.537	47.458	1.00 38.77	6
ATOM	205	С	PRO A	26		32.433	45.903	49.891	1.00 37.32	6
MOTA	206	0	PRO A			31.416	46.372	50.412	1.00 32.70	8
	207	N	ARG A			33.134	44.930	50.452	1.00 36.54	7
ATOM										
ATOM	208	CA	ARG A			32.685	44.359	51.711	1.00 37.39	6
MOTA	209	CB	ARG A			32.116	42.952	51.455	1.00 35.29	6
MOTA	210	CG	ARG A	27		31.047	42.956	50.355	1.00 32.69	6
ATOM	211	CD	ARG A	27		30.507	41.573	49.956	1.00 33.87	6
ATOM	212	NE	ARG A	27		29.757	40.909	51.021	1.00 36.16	7
ATOM	213	CZ	ARG A			30.293	40.132	51.959	1.00 37.11	6
ATOM	214		ARG A	27		31.604	39.903	51.976	1.00 34.42	7
ATOM	215		ARG A	27		29.516	39.597	52.896	1.00 33.67	7
ATOM	216	C	ARG A	27		33.813	44.329	52.732	1.00 36.35	6
MOTA	217	0	ARG A	27		33.881	45.188	53.610	1.00 35.77	8
ATOM	218	N	VAL A	28		34.703	43.351	52.607	1.00 34.93	7
MOTA	219	CA	VAL A	28		35.810	43.230	53.537	1.00 34.00	6
ATOM	220	CB	VAL A	28		36.633	41.954	53.252	1.00 36.21	6
ATOM	221		VAL A	28		37.574	41.652	54.424	1.00 33.59	6
ATOM	222		VAL A	28		35.696	40.790	52.992	1.00 37.05	6
ATOM	223	C	VAL A			36.712	44.454	53.423		
				28		_	44.959		1.00 31.91	6
ATOM	224	0	VAL A	28		37.216		54.427	1.00 31.45	8
MOTA	225	N	SER A	29		36.908	44.936	52.199	1.00 33.12	
ATOM	226	CA	SER A	29		37.751	46.111	51.967	1.00 32.03	6
MOTA	227	CB	SER A	29		38.205	46.181	50.499	1.00 31.77	6
MOTA	228	OG	SER A	29		37.113	46.223	49.600	1.00 30.80	8
MOTA	229	С	SER A	29		37.003	47.380	52.353	1.00 30.16	6
ATOM	230	0	SER A	29		37.604	48.404	52.650	1.00 28.70	8
ATOM	231	N	LEU A	30		35.682	47.310	52.352	1.00 32.43	7
MOTA	232	CA	LEU A	30		34.900	48:465	52.745	1.00 34.56	6
	233	CB	LEU A	30		33.463	48.358	52.221	1.00 36.44	6
ATOM .							49.513			
ATOM	234	CG	LEU A	30		32.508		52.560	1.00 36.79	6
ATOM	235		LEU A	30		32.070	49.446	54.012	1.00 36.73	6
MOTA	236		LEU A	30		33.202	50.840	52.256	1.00 37.84	6
MOTA	237	С	LEU A	30		34.902	48.527	54.262	1.00 34.89	6
MOTA	238	0	LEU A	30		35.033	49.601	54.841	1.00 37.58	8
ATOM	239	N	LEU A	31	•	34.761	47.366	54.897	1.00 34.07	7
ATOM	240	CA	LEU A	31		34.743	47.276	56.350	1.00 34.85	6
ATOM	241	CB	LEU A	31		34.768	45.808	56.791	1.00 36.37	6
ATOM	242	CG	LEU A	31		34.459	45.471	58.261	1.00 36.04	6
ATOM	243		LEU A	31		34.841	44.027	58.532	1.00 35.13	6
	244		LEU A			35.228	46.357	59.194	1.00 35.86	. 6
ATOM				31						
ATOM	245	C	LEU A	31		35.976	47.994	56.894	1.00 36.43	. 6
ATOM	246	0	LEU A	31		35.855	49.035	57.544	1.00 35.87	8
ATOM	247	N	LEU A	32		37.157	47.426	56.635	1.00 37.76	7
ATCM	248	CA	LEU A	32		38.420	48.015	57.087	1.00 36.82	б
ATOM	249	CB	LEU A	32		39.611	47.318	56.418	1.00 36.37	6
ATOM	250	CG	LEU A	32		40.030	45.888	56.774	1.00 39.11	6
ATOM	251		LEU A	32		41.117	45.420	55.815	1.00 35.16	6
ATOM	252		LEU A	32		40.538	45.830	58.214	1.00 37.73	6
	253					38.500	49.513	56.780	1.00 34.84	6
ATOM		C	LEU A	32						8
ATOM	254	0	LEU A	32		38.846	50.326	57.644	1.00 36.58	
ATCM	255	N	ARG A	33		38.184	49.877	55.545	1.00 31.37	7
ATOM	256	CA	ARG A	33		38.247	51.270	55.150	1.00 32.53	6
ATOM	257	CB	ARG A	33	٠.	37.927	51.398	53.662	1.00 31.52	6
ATCM	258	CG	ARG A	33		38.481	52.652	53.042	1.00 35.88	6
ATOM	259	CD	ARG A	33		38.107	52.752	51.581	1.00 43.44	6
ATCM	260	NE	ARG A	33			51.583	50.811	1.00 48.37	7
	261	CZ	ARG A	33		38.348	51.469	49.497	1.00 52.27	6
ATOM								48.823	1.00 52.27	7
ATCM	262		ARG A	33.		37.771	52.459			7
	263		ARG A	33		38.739	50.369	48.858	1.00 51.08	
ATCM	264	C	ARG A	33		37.274	52.102	_55.989 _.	1.00 32.32	, 6
				•	•					•

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MOTA	265	0	·ARG A	33		37.471	53.299	56.196	1.00 29.23	8
ATOM	266	N	PHE A	34		36.231	51.445	56.484	1.00 32.58	7
ATOM	267		PHE A			35.216	52.096	57.304	1.00 32.69	6
ATOM	268		PHE A			33.952	51.232	57.359	1.00 31.22	6
	269		PHE A	-		32.838	51.825	58,183	1.00 28.74	6
MOTA								57.700		
ATOM	270		1 PHE A			32.085	52.888		1.00 22.76	6
MOTA	271	CD:				32.551	51.322	59.456	1.00 28.09	6
MOTA	272	CE:	l PHE A	34		31.061	53.441	58.472	1.00 23.70	6
ATOM	273	CE	PHE A	34		31.524	51.873	60.235	1.00 24.59	6
ATOM	274	CZ	PHE A	34		30.781	52.929	59.741	1.00 21.39	6 -
ATOM	275	c	PHE A			35.734	52.319	58.719	1.00 33.45	6
ATOM	276	ō	PHE A	_ :		35.635	53.425	59.258	1.00 35.49	8
	277					36.276	51.264	59.323		7
ATOM		N	LYS A						1.00 34.52	
MOTA	278	CA	LYS A			36.805	51.360	60.678	1.00 36.51	6
MOTA	279	CB	LYS A			37.118	49.977	61.235	1.00 36.47	6
MOTA	280	CG	LYS A	35		35.912	49.074	61.343	1.00 40.81	6
ATOM	281	CD	LYS A	. 35		36.246	47.801	62.090	1.00 44.10	6
ATOM	282	CE	LYS A	35		37.347	47.029	61.402	1.00 47.46	6
ATOM	283	NZ	LYS A			38.601	47.823	61.276	1.00 53.53	7
ATOM	284	С	LYS A			38.054	52.222	60.735	1.00 36.61	6
MOTA	285	ŏ	LYS A			38.352	52.824	61.766	1.00 .36.78	8
	286	N	ASP A			38.794	52.267	59.635	1.00 36.27	7
MOTA										
MOTA	287	CA	ASP A			39.980	53.090	59.592	1.00 39.71	6
MOTA	288	CB	ASP A			40.679	52.937	58.239	1.00 44.78	6
MOTA	289	CG	ASP A			41.863	53.892	58.075	1.00 47.10	6
MOTA	290	OD1	. ASP A	36		42.803	53.852	58.906	1.00 44.02	8
MOTA	291	OD2	ASP A	36		41.843	54.682	57.106	1.00 48.43	8
ATOM	292	С	ASP A	36	•	39.508	54.530	59.789	1.00 39.99	6
ATOM	293	0	ASP A	36		40.023	55.258	60.636	1.00 40.76	8
ATOM	294	N	ALA A			38.506	54.919	59.007	1.00 38.59	7
ATOM	295	CA	ALA A	37		37.939	56.258	59.066	1.00 37.14	6
	296	CB	ALA A	37		36.857	56.402	58.000	1.00 35.85	6
MOTA							56.549			
ATOM	297	C	ALA A	37		37.354		60.446	1.00 38.34	6
ATOM	298	0	ALA A	37		37.391	57.687	60.928	1.00 37.32	8
ATOM	299	N	MET A	38		36.809	55.518	61.079	1.00 36.19	7
MOTA	300	CA	MET A	38		36.213	55.674	62.397	1.00 36.80	6
ATOM	301	CB	MET A	38		35.141	54.598	62.606	1.00 37.38	6
ATOM	302	.cg	MET A	38	-	33.938	54.717	61.673	1.00 37.60	6
ATOM	303	SD	MET A	38		32.887	56.165	61.999	1.00 33.61	16
ATOM .	304	CE	MET A	38		32.398	55.824	63.680	1.00 35.60	6
ATOM	305	C	MET A	38		37.262		63.502	1.00 35.84	6
ATOM	306	ō	MET A	38		36.937	55.688	64.692	1.00 34.89	8
ATOM	307	N	ASN A	39		38.518	55.400	63.100	1.00 33.83	7
	308	CA	ASN A	39		39.626	55.264	64.044	1.00 34.94	6
								64.775		6
MOTA	309	CB	ASN A	39		39.897	56.582		1.00 32.48	
MOTA	310	CG	ASN A	39 .		40.213	57.717	63.825	1.00 32.34	6
MOTA	311		ASN A	39		41.128	57.621	63.009	1.00 31.85	8
ATOM	312	ND2	ASN A	39		39.455	58.800	63.924	1.00 30.92	7
ATOM	313	С	ASN A	39		39.253	54.183	65.045	1.00 36.87	6
ATOM	314	0	ASN A	39		39.403	54.357	66.260	1.00 36.60°	8
ATOM	315	N	LEU A	40		38.752	53.067	64.518	1.00 37.48	7
ATOM	316	CA	LEU A	40		38.341	51.933	65.336	1.00 39.66	6
ATOM	317	CB	LEU A	40		36.863	51.622	65.086	1.00 41.35	5
	. 318	CG	LEU A	40		35.858	52.712	65.476	1.00 42.69	6
	319		LEU A			34.448	52.261	65.111	1.00 45.05	6
ATOM				40				66.966		6
ATOM	320		LEU A	40		35.951	52.989		1.00 39.44	
ATOM	321	С	LEU A	40		39.184	50.687	65.058	1.00 39.79	6
ATOM	322	0	LEU A	40		38.804	49.575	65.434	1.00 36.88	8
MOTA	323	N	ILE A	41		40.337	50.889	64.420	1.00 40.50	7
MOTA	324	CA	ILE A	41	-	41.237	49.790	64.068	1.00 41.39	6
ATOM	325	CB	ILE A	41		40.780	49.141	62.724	1.00 39.24	6
ATOM	326		ILE A	41		41.017	50.103	61.564	1.00 36.97	6
	327	CG1		41		41.513	47.824	62.482	1.00 36.76	6
ATOM			ILE A	41		41.085	46.715	63.403	1.00 35.59	6
ATOM	328					42.684		63.913	1.00 44.37	6
MOTA	329	C	ILE A	41			50.295			8
ATOM	330	0	ILE A	. 41		42.927	51,328	63.277	1.00 46.01	

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ATOM	331	N	ASP A	42		43.64	16	49.582	64.497	1.00	45.19	7
ATOM	332	CA	ASP A			45.04		49.982	64.372		45.62	6
MOTA	333	CB	ASP A	42		45.71	6	50.090	65.742		45.17	6
ATOM	334	CG	ASP A	42		44.96	6	51.005	66.682		44.43	6
ATOM	335	OD:	L ASP A	42		44.73	1	52.177	66.322		39.84	8
ATOM	336	OD	ASP A	42		44.61	.2	50.546	67.787		48.50	8
ATOM	337	C	ASP A	42		45.75	0	48.915	63.551		48.47	6
ATOM	338	0	ASP A	42		45.31	.6	47.757	63.547		49.85	8
ATOM	339	N	GLU A	. 43		46.83	0	49.288	62.864		49.24	7
ATOM	340	CA	GLU A	43		47.55	3	48.325	62.028		50.79	6
ATOM	341	CB	GLU A	43		48.82	0	48.956	61.431		49.90	6
MOTA	342	CG	GLU A	43		48.54	4	50.029	60.378		57.20	6
MOTA	343	CD	GLU A	43		49.80	8	50.537	59.690		59.56	6
ATOM	344	OE1				50.51	7	49.721	59.061		65.05	8
MOTA	345	OE2				50.09		51.750	59.772		57.82	8
ATOM	346	С	GLU · A			47.91	8	47.020	62.733	1.00	49.73	6
MOTA	347	0	GLU A			47.81		45.943	62.149	1.00	49.18	8
ATOM	348	N	LYS A	44		48.32		47.118	63.992	1.00	49.40	7
MOTA	349	CA	LYS A	44		48.73		45.949	64.762	1.00	49.09	6
ATOM	350	CB	LYS A	44		49.31		46.418	66.093		52.46	6
MOTA	351	CG	LYS A	44		50.44		47.421	65.899		55.75	6
ATOM	352	CD	LYS A	44		51.16		47.749	67.201		58.74	6
ATOM	353 354	CE	LYS A	44		52.32		48.704	66.949		58.48	6
ATOM ATOM	355	NZ C	LYS A	44		53.122		48.968	68.176		58.95	7
ATOM	356	Ö	LYS A	44 44		47.638		44.897	64.994		47.63	6
ATOM	357	N	GLU A	45		46.379		45.298	65.290	1.00		8
ATOM	358	CA	GLU A			45.268		44.374	64.854 65.046	1.00		7
ATOM	359	CB	GLU A	45	•	44.024		45.143	65.514		43.09 41.19	6 6
ATOM-	360	ÇG	GLU A	45		44.192		45.859	66.844	1.00		6
ATOM	361	CD	GLU A	45		43.003		46.741	67.204	1.00		6
ATOM	362	OE1	GLU A	45		42.707		47.701	66.447	1.00		8
ATOM	363	OE2	GLU A	45		42.368		46.479	68.253	1.00		8
ATOM	364	Ç	GLU A	45		44.965	9	43.660	63.726	1.00		6
ATOM .	365.	0	.GLU A	4.5		44.480	0	42.523	63.699	-1.00		8
ATOM	366	N	LEU A	46		45.282		44.341	62.632	1.00	40.29	7
ATOM	367	CA	LEU A	46		45.042		43.823	61.299	1.00	37.16	6
ATOM	368	CB	LEU A	46		44.910		44.990	60.331	1.00		6
ATOM	369	CG	LEU A	46		44.822		44.658	58.845	1.00		6
MOTA	370		LEU A	46		43.655		43.726	58.563	1.00		6
ATOM	371		LEU A	46		44.673		45.964	58.080	1.00		6
ATOM ATOM	372 373	С 0	LEU A	46 46		46.090 47.275		42.860	60.774	1.00		6
ATOM	374	N	ILE A	47		45.646		43.192 41.662	60.698	1.00		8
ATOM	375	CA	ILE A	47		46.540		40.657	60.406 59.844	1.00		7
ATOM	376	CB	ILE A	47		46.333		39.253	60.491	1.00		6 6
ATOM	377	CG2	ILE A	47		47.346		38.262	59.930	1.00		6
ATOM	378		ILE A	47		46.504		39.328	62.010	1.00		6
ATOM	379		ILE A	4.7		47.858		39.846	62.448	1.00		6
ATOM	380	С	ILE A	47		46.196		40.570	58.362	1.00		6
ATOM	381	0	ILE A	47		45.037		40.342	58.003	1.G0		8
ATOM	382	N	LYS A	48		47.194		40.772	57.504	1.00		7
ATOM	383	CA	LYS A	48	4	46.985	,	40.713	56.056	1.00		6
ATOM	384	CB	LYS A	48	4	48.258	}	41.087	55.308	1.00		6
MOTA	385	CG	LYS A	48	4	48.056	i	41.273	53.811	1.00	24.90	6
MOTA	386	CD	LYS A	48	4	49.389)	41.352	53.091	1.00	26.39	6
ATOM	387	CE	LYS A	48		49.233		41.864	51.679	1.00	27.71	6
MOTA	388	NZ	LYS A	48		48.774		43.275	51.696	1.00		7
MOTA	389	C	LYS A	48		46.595		39.299	55.654	1.00		6
atom	390	0	LYS A	48		47.072		38.325	56.235	1.00		8
ATOM	391	N	SER A	49		15.735		39.183	54.653	1.00		7
MOTE	392	CA	SER A	49		45.299		37.876	54.205	1.00		6
ATOM	393	CB	SER A	49		13.952		37.979	53.479	1.00		6
ATOM	394	OG C	SER A	49		12.911		38.329	54.373	1.00		8
ATOM	395	C	SER A	49 49		16.322		37.211	53.293	1.00		6
ATOM	396	0	SER A	77	4	17.095		37.885	52.612	1.00	JI.89	8

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ATOM	397	7 N	ARG	A	50		46.319	35.879	53.296	1.00 29.71	-
MOTA	398	3 C2					47.21				7 6
ATOM	399	CI			50		48.249		53.318	1.00 26.20	
MOTA	400) C			50		47.687		54.172		6
ATOM	401	. CI			50		48.818				6
ATOM.	402	. NE			50		48.359		55.762		. 6
MOTA	403				50		47.708		55.345	1.00 19.20	7
ATOM	404		11 ARG		50		47.430		54.055		6
ATOM	405		12 ARG		50		47.334				7
MOTA	406		ARG		50		46.370		51.723		7
MOTA	407		ARG		50		45.319		52.206	1.00 23.30	6
ATOM	408	N	PRO		51		46.823		50.534		8
ATOM	409				51		48.021		49.789		7
ATOM	410		-		51		46.086		49.761		6
ATOM	411				51		46.862		48.451		6
MOTA	412	CG			51		47.503		48.392		6
ATOM	413		PRO		51		46.153		50.498		6
ATOM	414	0	PRO		51		47.071		51.293	1.00 31.32	6
ATOM	415	N	ALA		52		45.176	30.435	50.250	1.00 26.02	8 7
ATOM ·	416	CA	ALA	Α	52	•	45.151		50.876	1.00 25.76	6
ATOM	417	CB	ALA	Α	52		43.720		50.933	1.00 21.42	6
ATOM	418	С	ALA	Α	52		46.013		50.000	1.00 26.31	6
ATOM	419	0	ALA	A	52		45.878		48.780	1.00 30.31	. 8
ATOM	420	N	THR	A	53		46.909		50.608	1.00 26.80	. 7
ATOM	421	CA	THR	A	53		47.759		49.831	1.00 27.52	6
MOTA	422	CB	THR	A	53		48.845	25.975	50.717	1.00 26.27	6
ATOM	423	OG.			53		48.255	25.053	51.641	1.00 29.51	8
MOTA	424	CG.			53		49.522	27.076	51.502	1.00 24.66	6
ATOM	425	С	THR		53		46.908	25.462	49.209	1.00 26.58	6
MOTA	426	0	THR		53		45.778	25.228	49.634	1.00 21.98	8
ATOM	427	N	LYS		54		47.455	24.782	48.203	1.00 29.62	7
MOTA	428	CA	LYS		54		46.739	23.713	47.507	1.00 32.62	6
MOTA	429	CB	LYS		54		47.601	23.151	46.370	1.00 31.99	6
ATOM	430	CG	LYS		54 .		46.985	21.967	45.629	1.00 36.62	6
ATOM	431	CD	LYS		54		45.733	22.352	44.866	1.00 40.69	6
ATOM ATOM	432 433	CE NZ	LYS		54		46.058	23.173	43.625		6
ATOM	434	C	LYS		54		46.844	22.393	42.614	1.00 50.68	7
ATOM	435	o	LYS		54 54		46.348	22.595	48.465	1.00 36.00	6
ATOM	436	N	GLU		55		45.277	21.991	48.330	1.00 34.77	8
ATOM	437	CA	GLU		55		47.216 46.979	22.336	49.443	1.00 37.91	7
ATOM	438	CB	GLU		55		48.240	21.290 21.100	50.433	1.00 36.96	6
ATOM	439	ĊĠ	GLU		55		48.216	19.887	51.281 52.195	1.00 40.29	6
ATOM	440	CD	GLU		55		49.552	19.654	52.891	1.00 47.95 1.00 51.01	6
ATOM	441	OË1			55		49.659	18.688	53.679	1.00 52.65	6 8
ATOM	442	OE2			55		50,497	20.437	52.646	1.00 52.65	8
ATOM	443	С	GLU .		55		45.771	21.609	51.322	1.00 34.10	6
ATOM -	444	0	GLU .	A	55	-	44.892	20.769	51.496	1.00 33.08	8
MOTA	445	N	GLU .	A	56		45.723	22.827	51.866	1.00 32.39	7
ATOM	446	CA	GLU .	A	56		44.621	23.256	52.733	1.00 30.13	6
MOTA	447	CB	GLU Z	A	56		44.824	24.714	53.177	1.00 25.28	6
ATOM	448	CG	GLU 3	Ą	56		46.204	24.994	53.758	1.00 28.82	6
MOTA	449	CD	GLU 3	A	56		46.421	26.450	54.181	1.00 30.74	6
MOTA	450	OE1	GLU 2	4	56		46.072	27.369	53.398	1.00 29.77	8
ATOM	451	OE2	GLU A	A	56		46.969	26.674	55.288	1.00 25.98	8
ATOM	452	C	GLU A	A	56		43.264	23.114	52.024	1.00 29.63	6
ATOM	453	0	GLU ?		56		42.299	22.584	52.595	1.00 29.90	8
MOTA	454	N	LEU A		57		43.188	23.581	50.780	1.00 26.76	7
ATOM	455	CA	LEU A	•	57		41.944	23.490	50.020	1.00 25.29	6
ATOM	456	CB	LEU A	A.	57		42.132	24.103	48.629	1.00 22.68	6
ATOM	457	CG	LEU A		57		42.402	25.612	48.572	1.00 22.39	5
MOTA	458	CD1	LEU A	٠.	57		42.654	26.045	47.123	1.00 20.77 .	6
ATOM	459		LEU A		57		41.211	26.366	49.156	1.00 17.66	6
ATCM	460	C	LEU A		57		41.479	22.037	49.896	1.00 26.02	6
	461		LEU A		57		40.284	21.741	50.014	1.00 23.41	8
ATOM	462	N	LEU A		58		42.444	21.143	49.675	1.00 24.82	7
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MOTA MOTA MOTA	463 464 465	CA LEU . CB LEU . CG LEU .	A 58	42.194 43.434 43.838	19.027	49.526 48.965 47.558	1.00 22.44 1.00 21.02 1.00 22.94	6 6
ATOM	466	CD1 LEU	A 58	45.212		47.176	1.00 20.35	6 6
ATOM	467	CD2 LEU		42.755		46.587	1.00 23.28	6
MOTA MOTA	468 469	C LEU		41.797		50.835	1.00 25.20	6
MOTA	470	N LEU		41.456 41.858	17.867 19.794	50.854	1.00 26.55	8
ATOM	471	CA LEU	-	41.446	19.794	51.938 53.211	1.00 25.44	7
ATOM	472	CB LEU		41.559	20.229	54.350	1.00 25.24 1.00 24.68	6 6
ATOM	473	CG LEU A		42.956	20.490	54.912	1.00 27.05	6
MOTA	474	CD1 LEU 2		42.912	21.565	56.001	1.00 24.76	6
MOTA MOTA	475 - 476	CD2 LEU 2		43.492	19.184	55.474	1.00 26.99	6
ATOM	477	O LEU A		39.991 39.548	18.807 17.794	53.045 53.581	1.00 24.22	6
ATOM	478	N PHE	_	39.270	19.615	52.270	1.00 21.18	8 7
MOTA	479	CA PHE A		37.859	19.403	52.011	1.00 25.00	6
ATOM	480	CB PHE A		37.054	20.560	52.605	1.00 26.34	6
ATOM ATOM	481 482	CG PHE A		35.600	20.555	52.223	1.00 29.37	6
ATOM	483	CD2 PHE A		34.811 35.015	19.422 21.692	52.427 51.661	1.00 27.57	6
ATOM	484	CE1 PHE A		33.466	19.419	52.077	1.00 27.33 1.00 27.00	6 6
ATOM	485	CE2 PHE A		33.670	21.699	51.306	1.00 28.08	6
MOTA	486	CZ PHE A		32.893	20.559	51.513	1.00 29.48	6
ATOM ATOM	487 488	C PHE A		37.506 37.022	19.214	50.538	1.00 27.78	6
ATOM	489	N HIS A		37.734	18.143 20.220	50.158 49.696	1.00 31.57 1.00 26.76	. 8 7
MOTA	490	CA HIS A		37.376	20.056	48.287	1.00 28.84	6
ATOM	491	CB HIS A		37.365	21.405	47.561	1.00 27.76	6
MOTA MOTA	492 493	CC HIS A		36.385 35.056	22.396 22.549	48.117	1.00 30.54	6
ATOM	494	ND1 HIS A		36.750	23.401	47.907 48.987	1.00 33.74	6 7
MOTA	495	CE1 HIS A		35.691	24.135	49.286	1.00 32.07	6
ATOM	496	NE2 HIS A	61	34.649	23.638	48.644	1.00 34.10	7
ATOM ATOM	497 498	C HIS A O HIS A	61 61	38.278 39.287	19.056	47.539	1.00 28.38	6
ATOM	499	N THR A	62	37.895	18.604 18.705	48.072 46.310	1.00 25.81 1.00 32.88	8 7
ATOM	500	CA THR A	62	38.658	17.749	45.488	1.00 34.68	6
MOTA	501	CB THR A	62	37.715	16.739	44.778	1.00 34.36	6
MOTA MOTA	502 503	OG1 THR A	62 62	36.942 36.759	17.415 16.112	43.778	1.00 34.81	8
ATOM	504	C THR A	62	39.485	18.454	45.778 44.408	1.00 34.33 1.00 35.60	6 6
MOTA	505	O THR A	62	39.017	19.418	43.790	1.00 30.85	8
ATOM	506	N GLU A	63	40.700	17.958	44.166	1.00 37.38	7
ATOM ATOM	507 508	CA GLU A	63 63	41.587 42.759	18.555	43.165	1.00 40.68	6
ATOM		CG GLU A	63	43.719	17.626 17.389	42.840 43.987	1.00 43.75 1.00 50.68	6 6
ATOM		CD GLU A	63	45.026	16.760	43.529	1.00 55.36	6
ATOM		OE1 GLU A	63	45.789	17.441	42.808	1.00 53.03	8
ATOM ATOM	512 513	OE2 GLU A	63 63	45.285	15.585	43.883	1.00 59.56	8
ATOM		O GLU A	63	40.894 40.771	18.939 20.116	41.860 41.535	1.00 39.26	6
ATOM		N ASP A	64	40.453	17.948	41.102	1.00 42.33	8 7
MOTA		CA ASP A	64	39.782	18.224	39.845	1.00 36.98	6
ATOM		CB ASP A	64	38.957	17.000	39.426	1.00 42.19	6
ATOM ATOM		CG ASP A OD1 ASP A	64 64	38.037 37.039	16.501	40.533	1.00 47.66	6
ATOM		OD2 ASP A	64	38.325	17.193 15.413	40.851	1.00 47.95 1.00 50.07	8
MOTA	521	C ASP A	64	38.908	19.480	39.906	1.00 33.40	6
MOTA		O ASP A	64	38.927	20.293	38.986	1.00 33.64	8
ATOM		N TYR A	65 65	38.156	19.641	40.990	1.00 30.57	7
· ATOM ATOM		CA TYR A	65 65	37.286 36.300	20.806	41.157 42.316	1.00 29.65 1.00 30.16	6 6
ATOM		CG TYR A	65	35.557	21.790	42.316	1.00 30.16	6
ATOM	527	CD1 TYR A	65	34.791	22.572	41.944	1.00 30.25	6
ATOM	528	CE1 TYR A	65	34.126	23.715	42.399	1.00 28.36	6
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ATOM	529 CD2 TYR A 65	35.63	8 22.181 44	.150 1.00 28 28 6
ATOM	530 CE2 TYR A 65	34.98		
ATOM	531 CZ TYR A 65			.617 1.00 26.96 6
ATOM		34.22		.740 1.00 29.79 6
		33.56		.207 1.00 28.53 8
MOTA	533 C TYR A 65	38.118	3 22.061 41.	418 1.00 29.15 6
MOTA	534 O TYR A 65	37.860		
ATOM	535 N ILE A 66	39.122		.857 1.00 30.45 8 .270 1.00 26.61 7
ATOM	536 CA ILE A 66	39.986		
ATOM				.597 1.00 26.35 6
		40.998		.687 1.00 26.25 6
ATOM	538 CG2 ILE A 66	42.009	23.753 43.	869 1.00 21.20 6
MOTA	539 CG1 ILE A 66	40.264	22.341 44	000 1 00
ATOM	540 CD1 ILE A 66	39.478		
ATOM	541 C ILE A 66	40.761		201
ATOM	542 O ILE A 66			381 1.00 28.07 6
ATOM		41.039		225 1.00 31.26 8
		41.125		521 1.00 28.47 7
ATOM	544 CA ASN A 67	41.902	22.898 39.	337 1.00 30.15 6
MOTA	545 CB ASN A 67	42.563		
MOTA	546 CG ASN A 67	43.712		FD0
ATOM	547 OD1 ASN A 67	44.674		
ATOM	548 ND2 ASN A 67			878 1.00 43.34 8
ATOM		43.626		956 1.00 37.14 7
		41.020	23.554 38.	314 1.00 28.41 6
MOTA	550 O ASN A 67	41.494	24.354 37.	
ATOM	551 N THRA 68	39.733	23.221 38.	
ATOM	552 CA THR A 68	38.787	23.791 37.	
ATOM	553 CB THR A 68	37.438	23.111 37.	
ATOM	554 OG1 THR A 68	37.620		250
ATOM	555 CG2 THR A 68		21.695 37.	
ATOM		36.549	23.591 36.	
		38.633	25.263 37.	732 1.00 22.13 6
MOTA	557 O THR A 68	38.529	26.088 36.	830 1.00 21.97 8
ATOM	558 N LEUA 69	38.645	25.582 39.	
ATOM	559 CA LEU A 69	38.535	26.956 39.	
ATOM	560 CB LEU A 69	38.376	26.982 41.	482 1.00 23.97 6
ATOM	561 CG LEU A 69	37.023		
MOTA	562 CD1 LEU A 69		26.527 41.9	
ATOM		37.087	26.416 43.0	066 1.00 30.99 6
	***	35.942	27.528 41.	120 1.00 28.69 6
MOTA	564 C LEU A 69	39.772	27.757 39.0	
ATOM	565 0 LEUA 69	39.683	28.921 38.6	
ATOM	566 N MET A 70	40.932	27.128 39.2	
MOTA	567 CA MET A 70	42.183	27.794 38.8	
ATOM	568 CB MET A 70	43.358		
ATOM	569 CG MET A 70	_	26.953 39.3	
ATOM		43.418	26.751 40.8	
		44.970	25.929 41.3	325 1.00 30.71 16
ATOM	571 CE MET A 70	46.137	27.077 40.6	
ATOM	572 C MET A 70	42.324	28.040 37.4	
ATOM	573 O MET A 70	42.903	29.041 36.9	
MOTA	574 N GLU A 71	41.769	27.122 3€ €	
ATOM	575 CA GLU A 71	41.859		
ATOM			27.204 35.1	
ATOM		41.681	25.814 34 5	
		42.224	25.695 33.1	67 1.00 31.75 6
ATOM	578 CD GLU A 71	43.737	25.905 33.0	
ATOM.	579 OE1 GLU A 71	44.288	25.855 31.9	83 1.00 35.84 8
ATOM	580 OE2 GLU A 71	44.377	26.116 34.1	
ATOM	581 C GLU A 71	40.845		
ATOM	582 O GLU A 71			
		41.144	28.851 33.6	
ATOM	583 N ALA A 72	39.649	28.197 35.1	69 1.00 19.22 7
ATOM	584 CA ALA A 72	38.589	29.067 34.6	84 1.00 19.39 6
MOTA	585 CB ALA A 72	37.298	28.743 35.3	97 1.00 19.23 6
ATOM	586 C ALA A 72	38.931	30.536 34.8	
MOTA	587 O ALA A 72	38.711		
MOTA	588 N GLU A 73			
		39.470	30.835 36.0	
ATOM	589 CA GLU A 73	39.820	32.202 36.4	
ATOM	590 CB GLU A 73	40.157	32.282 37.9	
ATOM	591 CG GLU A 73	40.646	33.655 38.3	
ATOM .	592 CD GLU A 73	40.840	33.806 39.8	
ATOM	593 OE1 GLU A 73	39.841	33.776 40.5	0 1 00 20 40 0
ATOM	594 OE2 GLU A 73	41.996		
ATOM		31.330	33.960 40.2	77 1.00 31.77 8
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ATOM	595	С	GLU	JA	73	40.94	6 32.840	35.615	1.00 31.83	6
ATOM	596	O	GLU		73	40.85			1.00 33.52	8
ATOM	597	N	ARG		74	41.99			1.00 31.45	7
ATOM	598	CA	ARG		74	43.12			1.00 30.65	6
ATOM	599	CB	ARG		74	44.40			1.00 32.12	6
ATOM	600	CG	ARG		74	44.51			1.00 32.12	
ATOM	601	CD	ARG		74	45.70				6
MOTA	602	NE	ARG		74	46.04			1.00 30.73	6
	603	CZ	ARG		74				1.00 34.18	7
ATOM						46.64			1.00 35.55	6
ATOM	604		L ARG		74	46.98			1.00 29.64	7
ATOM	605	NH2			74	. 46.90			1.00 34.07	7
ATOM	606	C	ARG		74	42.89		33.051	1.00 28.61	6
ATOM .	607	0	ARG		74	43.43		32.338	1.00 24.38	8
MOTA	608	N	CYS		75	42.10		32.566	1.00 28.32	7
ATOM	609	CA	CYS		75	41.79		31.148	1.00 32.42	6
ATOM	610	CB	CYS		75	41.68		30.682	1.00 32.91	6
MOTA	611	SG	CYS		75	43.28		30.777	1.00 37.09	16
ATOM	612	С	CYS		75	40.48		30.956	1.00 33.10	
MOTA	613	0	CYS		75	40.02		29.834	1.00 30.74	8
MOTA	614	N	GLN		76	39.91		32.088	1.00 34.42	7
ATOM -	61,5	CA	GLN		76	38.69		32.144	1.00 33.20	6
MOTA	616	CB	GLN		76	38.98		31.578	1.00 32.09	5
MOTA	617	CG	GLN		76	38.08		32.094	1.00 39.46	6
MOTA	618	CD	GLN	A	76	38.47	9 36.541	33.480	1.00 41.47	6
ATOM	619	OE1			76	38.57	4 35.755	34.426	1.00 45.02	`8
MOTA	620	NE2			76	38.70		33.606	1.00 42.22	7
MOTA	621	С	GLN		76	37.56		31.358	1.00 33.20	6
ATOM	622	0	GLN		76	. 36.73	2 33.598	30.760	1.00 34.19	8
MOTA	623	N	CX2		7 7	37.52		31.370	1.00 31.81	7
MOTA	624	CA	CYS		77	36.51	1 30.862	30.627	1.00 31.47	6
MOTA	625	CB	CYS	$\cdot A$	77	37.18	7 30.181	29.454	1.00 30.25	6
ATOM	626	SG	CYS	А	77.	38.47	9 29.071	30.044	1.00 33.94	16
MOTA	627	С	CYS	Α	77	35.85	1 29.795	31.498	1.00 31.97	6
MOTA	628	Ö	CYS		77	36.33	5 29.503	32.590	1.00 35.15	8
ATOM	629	N	VAL	A	78	34.75	0 29.216	31.018	1.00 30.78	7
ATOM	630	CA	VAL	A	78	34.06	9 28.139	31.747	1.00 30.55	6
ATOM	631	CB	VAL	A	78	32.53	9 28.287	31.720	1.00 30.06	6
MOTA	632		VAL		78	31.88		32.293	1.00 28.23	б
MOTA	633		VAL		78	32.12		32.526	1.00 30.67	6
ATOM	634	С	VAL		78	34.42	0 26.794	31.110	1.00 29.80	6
MOTA	635	0	VAL		78	33.85	1 26.422	30.077	1.00 29.65	8
MOTA	636	Ŋ	PRO		79	35.33	7 26.033	31.739	1.00 28.55	7
MOTA	637	CD	PRO		79	35.98	5 26.335	33.025	1.00 24.39	6
ATOM	638	CA	PRO		79	35.79	3 24.724	31.261	1.00 28.89	6
ATOM	639	CB	PRO		79	36.62	2 24.218	32.434	1.00 24.49	6
MOTA	6.0	CG	PRO	A	79	37.23	9 25.500	32.922	1.00 25.68	6
MOTA	61	С	PRO	A	79	34.66	8 23.776	30.881	1.00 30.13	6
MOTA	6.2	0	PRO	A	79	33.69	7 23.624	31.615	1.00 30.87	8
MOTA	643	N	LYS	A	30	34.79	5 23.136	29.727	1.00 33.44	7
MOTA	644	CA	LYS	A	80	33.75	3 22.216	29.303	1.00 38.52	6
MOTA	645	CB	LYS	A	80	34.20	2 21.421	28.076	1.00 45.18	6
ATOM	646	CG	LYS	A	80	35.45	20.589	28.278	1.00 55.18	6
MOTA	647	CD	LYS	A	80	35.78	3 19.827	27.000	1.00 60.80	6
ATOM	. 648	CE	LYS		80	37.03	5 18.976 [.]	27.168	1.00 64.25	6
ATOM	649	NZ	LYS	A	80	37.36		25.911	1.00 68.95	7
ATOM	650	С	LYS		80	33.41		30.443	1.00 36.56	6
ATOM	651	0	LYS		80	34.29		31.164	1.00 31.61	8
ATOM	652	N	GLY		81	32,11		30.602	1.00 32.57	7
ATOM	653	CA	GLY		81	31.63		31.648	1.00 29.81	6
ATOM	654	C	GLY		81	31.47		32.965	1.00 28.30	6
ATOM	655	õ	GLY		81	30.54		33.723	1.00 25.49	8
ATOM	656	N	ALA		82	32.38		33.218	1.00 25.99	7
ATOM	657	CA	ALA		82	32.38		34.458	1.00 26.72	6
	658	CB	ALA		82	33.48		34.406	1.00 22.64	6
ATOM	659	C	ALA		82	31.06		34.886	1.00 27.84	6
ATOM		0	ALA		82	30.72		36.068	1.00 30.00	8
ATOM	660	J	- ALL	<u> </u>	02	30.72	23.224	20.008	1,00 30.00	U

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•						
ATOM	661 N ARG A 83	30.31	0 22 01.			
MOTA	CC0 01	_				7
MOTA	CC2 CD	29.07		2 34.349	5 1.00 32.50	6
		28.28	5 24.94:	1 33.127	7 1.00 37.19	
ATOM	664 CG ARG A 83	27.439	26.189		2.00 37.13	6
MOTA	665 CD ARG A 83	26.480				6
ATOM-	666 NE ARG A 83	25.904				6
ATOM	CC7 OF 170					7
ATOM	CCO 1774 177	25.046			1.00 56.84	6
		24.649	26.413	36.724	1.00 53.05	
MOTE	669 NH2 ARG A. 83	24.588	28.672			7
ATOM	670 C -ARG A 83	28.208				. 7
ATOM	671 O ARG A 83					6
ATOM	(77) 11	28.056			1.00 29.62	8
	C72 02	27.648		34.581	1.00 33.06	7
MOTA	673 CA GLUA 94	26.819		35.343		
ATOM	-674 CB GLUA 84	26.112	20.562		1.00 35.40	. 6
ATOM	675 CG GLU A 84	26.989				· 6
ATOM	676 CD GLU A 84					6
ATOM	C77 074	27.551				6
	6// OE1 GLU A 84	27.925	19.723	31.292	1.00 41.12	
MOTA	678 OE2 GLU A 84	27.636	21.671	32.270		8
MOTA	679 C GLUA 84 ·	27.617	20.823	36.417		8
ATOM	680 O GLUA 84	27.246		20.41/		6
ATOM	681 N LYS A 85		20.816		1.00 34.66	8
ATOM	C00	28.727	20.226	36.002	1.00 35.21	7
	CO2 CD	29.604	19.450	36.878	1.00 37.93	6
ATOM	683 CB LYS A 85	30.841	19.030	36.076	1.00 40.61	
ATOM	684 CG LYS A 85	31.739	17.977	36.706	1 00 40.61	6
MOTA	685 CD LYS A 85	31.038	16.640		1.00 42.63	6
ATOM	686 CE LYS A 85			36.872	1.00 45.48	6
ATOM	COS	32.054	15.523	37.078	1.00 45.60	6
ATOM	COO	33.032	15.833	38.154	1.00 46.16	7
		30.032	20.159	38.175	1.00 37.56	
ATOM	689 O LYS A 85	30.161	19.516	39.222	1.00 38.40	6
MOTA	690 N TYRA 86	30.254	21.472	38.116	1.00 38.40	8
ATOM	691 CA TYR A 86	30.671			1.00 35.60	7
ATOM	692 CB TYR A 86		22.216	39.307	1.00 32.67	б
ATOM	602	32.151	22.610	39.200	1.00 32.09	6
		33.065	21.424	38.995	1.00 33.63	6
MOTA	694 CD1 TYR A 86	33.120	20.393	39.932	1.00 32.12	
ATOM	695 CE1 TYR A 86	33.918	19.266	39.723	1 00 32 50	6
MOTA	696 CD2 TYR A 86	33.839	21.306		1.00 33.59	6
ATOM	697 CE2 TYR A 86	34.645		37.841	1.00 33.82	6
ATOM	500 00		20.178	37.623	1.00 34.55	6
ATOM	£00	34.675	19.162	38.566	1.00 32.38	6
	700	35.431	18.034	38.336	1.00 29.17	8
ATOM	700 C TYR A 86	29.831	23.455	39.597	1.00 30.21	
ATOM	701 O TYRA 86	30.192	24.265	40.445		6
ATOM	702 n asna 87	28.712	23.594		1.00 29.12	8
MOTA	703 CA ASN A 87			38.893	1.00 29.44	7
ATOM	704	27.797	24.717	39.086	1.00 28.58	6
TOM	705	27.154	24.618	40.470	1.00 25.63	6
		25.871	25.428	40.596	1.00 28.05	6
HOM	706 OD1 ASN A 87	25.275	25.477	41.672	1.00 11.32	
TOM	707 ND2 ASN A 87	25.434	26.055	39.506		8
-TOM	708 C ASN A 87	28.580	26.015		1.00 .8.35	7
MOTA	709 O ASN A 87			38.963	1.00 : 3.35	6
TOM	710	28.319	26.981	39.677	1.00 32.07	. 8
	711	29.545	26.019	38.051	1.00 32.05	· 7
MOT	711 CA ILE A 88	30.407	27.173	37.809	1.00 33.77	6
MOT	712 CB ILE A 88	31.894	26.734	37.776	1.00 36.13	
TOM	713 CG2 ILE A 88		27.831	37.201		6
TOM	714 CG1 TLE A 88	32.357	27.031		1.00 37.80	6
TOM	715 CD1 ILE A 88	32.357	26.342	39.178	1.00 38.92	6
	***************************************		27.483	40.176	1.00 41.44	6
TOM	716 C ILE A 88	30.085	27.857	36.482	1.00 32.28	6
TCM	717 O ILE A 88	29.708	27.196	35.520	1.00 32.72	
TOM.	718 N GLY A 89		29.179		1 00 34 7 7	8
TOM	719 CA GLY A 89			36.438	1.00 31.56	7
TOM			29.915	35.207	1.00 30.84	6
			30.689	35.093	1.00 32.17	6
TOM	721 O GLY A 89	28.628	31.670	34.349	1.00 30.42	8
TOM	722 N GLY A 90		30.257	35.821.	1.00 31.51	
TOM	723 CA GLY A 90				1 00 32.31	7
TOM ·	724 C GLY A . 90		30.53/	35.756	1.00 32.92	5
TCM				36.524	1.00 34.32	6
-				37.129	1.00 33.97	8
TOM	726 N TYR A 91	25.144	32.882	36.504	1.00 33.88	7
	• • •	•	-		2.00	

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MOTA	727	CA TYR	RA 91	24.92	4 34.146	37.206	1 00 25 76	_
MOTA	728	CB TYR		23.46				6
ATOM	729			23.08				6
ATOM	730				_	37.990		6
				23.41		37.688		6
MOTA	731			23.10	5 38.106	38.577	1.00 42.44	5
MOTA	732		A 91	22.44	4 35.484	39.205		
ATOM	733	CE2 TYR	A 91	22.13		40.097		6
MOTA	734			22.46		30.037		6
ATOM	735					39.775		6
				22.13	0 38.835	40.646	1.00 43.69	8
MOTA	736	C TYR		25.24	2 34.082	38.701	1.00 34.15	6
MOTA	737	O TYR	A 91	25.82	1 35.014	39.266	1.00 29.52	8
ATOM	738	N GLU	A 92	24.83		39.333	1.00 34.78	
ATOM	. 739	CA GLU		25.02				7
MOTA	740	CB GLU				40.767	1.00 38.46	6
ATOM	741			24.23		41.211	1.00 43.99	6
				23.932		42.700	1.00 52.10	. 6
ATOM	742	CD GLU		23.29	30.161	43.097	1.00 58.00	6
MOTA	743	OE1 GLU	A 92	24.001	29.126	43.058	1.00 60.63	8
ATOM	744	OE2 GLU	A 92	. 22.087		43.434	1.00 59.58	
ATOM	745	C GLU	A 92	26.492		41.208		8
ATOM	746	O GLU		26.902			1.00 36.42	5
ATOM	747	N ASN				42.193	1.00 32.92	8
				27.280		40.473	1.00 34.12	7
ATCM	748	CA ASN		28.693	31.671	40.808	1.00 33.24	6
MOTA	749	CB ASN	A 93	28.871	30.259	41.364	1.00 28.52	6
ATOM .	750	CG ASN	A 93	27.734		42.299	1.00 27.45	
ATOM	751	OD1 ASN	A 93	27.547		43.355		6
MOTA	752	ND2 ASN		26.956			1.00 21.76	8
ATOM	753	C ASN				41.895	1.00 21.79	7
MOTA	754			29.529		39.535	1.00 35.04	6
				30.160		39.059	1.00 33.81	8
ATOM	755	N PRO		29.583		39.01 0	1.00 36.19	7
MOTA	756	CD PRO	A 94	28.970	34.231	39.690	1.00 34.62	6
ATOM	757	CA PRO	A 94	30.274	33.560	37.808	1.00 34.80	
ATOM	758	CB PRO	A 94	29.924		37.791		6
ATOM	759	CG PRO		28.619			1.00 33.94	6
ATOM	760	C PRO			35.095	38.516	1.00 36.13	6
	761			31.775	33.379	37.733	1.00 34.63	6
ATOM		O PRO		32.443	33.103	38.730	1.00 34.72	8
ATOM	762	N VAL	. –	32.299	33.556	36.526	1.00 33.57	7
ATOM	763	CA VAL		33.735	33.499	36.307	1.00 30.31	6
ATOM	764	CB VAL	A 95	34.085	33.171	34.841	1.00 29.88	6
ATOM	765	CG1 VAL 2	A 95	35.561	33.453	34.574	1.00 29.53	
ATOM	766	CG2 VAL		33.795	31.713		1.00 29.55	6
ATOM	767	C VAL				34.563	1.00 28.05	6
ATOM	768	O VAL		34.195	34.910	36.624	1.00 29.86	6
				33.524	35.879	36.272	1.00 29.07	8
MOTA	769	N SER		35.318	35.019	37.317	1.00 30.89	7
ATOM	770	CA SER A		35.889	36.310	37.687	1.00 32.27	6
ATOM	771	CB SER A	4 96	34.885	37.145	38.501	1.00 30.16	6
MOTA	772	OG SER 3	A 96	34.600	36.545	39.756	1.00 26.77	8
MOTA	773	C SER A	96	37.111	35.993	38.537		
MOTA	774	O SER 3	96	37.603			1.00 32.96	6
ATOM	775	N TYR A	97		34.865	38.511	1.00 33.77	8
				37.609	36.973	39.282	1.00 32.66	7
ATOM	776	CA TYR A		38.753	36.712	40.132	1.00 31.95	6
MOTA	777	CB TYR A	97	39.838	37.766	39.923	1.00 31.81	6
ATOM	778	CG TYR A	97	40.416	37.729	38.525	1.00 30.39	6
MOTA	779	CD1 TYR A		39.820	38.434	37.479		
ATOM	780	CE1 TYR A		40.327			1.00 30.63	6
	781	CD2 TYR A			38.358	36.178	1.00 28.49	6
ATOM				41.536	36.945	38.236	1.00 28.43	6
ATOM		CE2 TYR A		42.046	36.858	36.942	1.00 24.73	6
ATOM	783	CZ TYR A		41.437	37.565	35.919	1.00 27.27	6
atom	784	OH TYR A	97	41.915	37.455	34.633	1.00 26.70	
ATOM		C TYR A		38.350			1 00 21 10	8
					36.618	41.596	1.00 31.10	6
ATOM				39.178	36.735	42.495	1.00 33.01	8
atom		N ALA A		37.059	36.398	41.818	1.00 31.11	7
ATOM		CA ALA A		36.510	36.241	43.160	1.00 30.06	5
ATOM	789	CB ALA A	98	35.141	36.920	43.256	1.00 27.71	6
ATCM	790	C ALA A	98	36.350	34.736	43.357	1.00 31.24	6
ATCM		O ALA A	98	36.335				
		N MET A	99		34.238	44.487	1.00 29.66	-8
MOTA	, 34	· HELL	ر ر	36.249	34.030	42.230	1.00 29.50	7

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ATO:	M 793 CA MET A 99	
ATO		36.048 32.589 42.207 1.00 29.89 6
	M 705 00 10 10 10	35.774 32.123 40 778 1.00 23.03 6
ATO		3E 040 30 55
ATO	A 33	36 426 31 031 032 1.00 29.63 6
OTA	:::: n jj	3E COD 15 30.120 1.00 49.78 16
OTA	M 798 C MET A 99	
OTA	M 799 O MET A 99	31.800 42.783 1.00 30 81 6
ATO	M 800 N PHE A 100	30./5/ 43.406 1.00 30 50
ATON	1112 A 100	38.417 32.274 42.569 1 00 32 00 7
ATON	1111 N TOO	39.554 31 557 42 114
		40 222 24 24
ATOM		41 424 20 25 1.00 33.95 6
ATOM	4 804 CD1 PHE A 100	47 152 22 50.070 41.14 6
ATOM	805 CD2 PHE A 100	42 760 20 222 43.364 1.00 41.84 6
ATOM	806 CE1 PHE A 100	42.372 1.00 42.18 6
ATOM	807 CE2 PHE A 100	28.115 43.941 1.00 43 63 6
ATOM	808 CZ PHE A 100	43.000 43.000 47 944 1 00 40
ATOM	IME A 100	43.517 28.487 43 720 1 00
ATOM	- 122 K 100	40.519 32.438 43 895 1 00 33.63
	- IMB A 100	
ATOM	2444 74 101	41.137 33 415 43 245 1.00 38.21 8
ATOM	m. n 101	42 002 2
MOTA	813 CB THR A 101	42 622 25 43.503 1.00 22.19 6
ATOM	814 OG1 THR A 101	43.072 1.00 22.48 6
ATOM	815 CG2 THR A 101	34.795 42.052 1.00 21 99 9
MOTA	816 C THR A 101	30.335 43.876 1.00 15 00
ATOM	817 O THR A 101	41.408 34.860 45.205 1 00 21 77
ATOM		41.988 34.845 46 282 1 00 22
ATOM		******* 33.3// 45 NEB 1 00 22
	819 CA GLY A 102	39.533 35.947 46 231 1 00 21.79 7
ATOM	820 C GLY A 102	30 070 21.23 6
ATOM	821 O GLY A 102	20 200
ATOM	822 N SER A 103	20 510 4
ATOM	823 CA SER A 103	30 000 7
ATOM	824 CB SER A 103	37 454 32 276 1.00 26.51 6
ATOM	825 OG SER A 103	36.334 31.398 46.314 1.00 28.10 6
ATOM	826 C SER A 103	32.099, 45.639 1.00 32 01 0
ATOM	827 O SER A 103	32.040 48.032 1.00 27 73 6
ATOM		31.544 49.144 1 00 30 63 0
ATOM		40.364 32.080 47.410 1.20.00
ATOM	829 CA SER A 104	
	830 CB SER A 104	42 760 24 40.33 6
MOTA	831 OG SER A 104	42 501 22 6
ATOM	832 C SER A 104	A1 070 70 400
ATOM	833 O SER A 104	42 026 21 22 22 2.00 23.57 6
ATOM	834 N LEU A 105	41 000 70 8
ATOM	835 CA LEU A 105	42 162 23.91 7
MOTA	836 CB LEU A 105	42 040 25 11 1
ATOM	837 CG LEU A 105	42.049 36.082 49.382 1.00 23 57 6
MOTA	838 CD1 LEU A 105	37.091 49.672 1.00 26 30 6
ATOM	839 CD2 LEU A 105	44.502 36.551 49.178 1.00 22.38 6
ATCM		32.023 38.413 48.984 1.00 27 36 6
MOTA		41.187 34.559 51.182 1.00 23 49 6
ATOM		41.604 34.448 52.331 1.00 27.50
	242	39.887 34.556 50 897 1 00 25 20
ATOM	843 CA ALA A 106	38.884 34 423 51 057 3 30 33.32
ATOM	844 CB ALA A 106	37.471 34 423 51 350 1 00 20.04 6
ATOM	845 C ALA A 106	20 000 25 25 25 25 26 24 28 8
ATOM.	846 O ALA A 106	30 053
ATOM	847 N THR A 107	20 4:0 2.00 42./2 8
MOTA	848 CA THR A 107	30.410 32.057 52.111 1.00 25.65 7
ATOM	849 CS THR A 107	33.020 30.760 52.754 1.00 25.54 6
ATOM	850 OG1 THR A 107	33.708 29.637 51.713 1.00 21.92 6
ATOM	851 CG2 THR A 107	20.33 49.688 50.868 1.00 26 40 0
ATOM	852 C THR A 107	39.742 28.295 52.387 1.00 17 36 6
ATOM		40.901 30.720 53 583 1 00 33 56
	853 O THR A 107	40.906 30.254 54.727 1.00 28.16 6
ATOM	854 N GLY A 108	44 00.
ATOM	855 CA GLY A 108	42 24
ATOM ·	856 C GLY A 108	47 00- 1-1-1 33:710 4:00 48:37 6
ATOM	857 O GLY A 108	42 500 30.20
ATOM	858 N SER A 109	43.302 31.499 56.076 1.00 32.98 8
		42.283 33.018 54.942 1.00 24.81 7

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Figure 17-14

MOTA	859	CA SER A 109	42.002	33.810	56.119	1.00 24.86	-
MOTA	860		41.222				6
ATOM	861				55.727	1.00 24.74	6
			41.992		54.872	1.00 21.07	8
ATOM	862		41.240	32.996	57.173	1.00 27.89	6
MOTA	863	O SER A 109	41.424	33.214	58.377	1.00 30.92	8
ATOM	864		40.389		56.744	1.00 23.91	7
ATOM	865		39.676				
					57.721	1.00 24.80	6
ATOM	866		38.641	30.290	57.074	1.00 29.65	6
ATOM	867		37.469	31.016	56.669	1.00 30.45	8
MOTA	868	CG2 THR A 110	38.228		58.067	1.00 29.00	6
ATOM	869		40.712				
ATOM	870				58.478	1.00 24.34	6
			40.615	30.282	59.699	1.00 24.74	8
ATOM	871		41.715	29.954	57.764	1.00 23.01	7
ATOM	872	CA VAL A 111	42.759	29.173	58.416	1.00 24.13	6
ATOM	873	CB VAL A 111	43.695	28.495	57.391	1.00 25.77	6
ATOM	874		44.845	27.773			
ATOM	875				58.121	1.00 22.51	6
			42.888	27.502	56.534	1.00 22.67	6
ATOM	876		43.576	30.071	59.329	1.00 23.14	6
MOTA	877	O VAL A 111 .	43.720	29.793	60.518	1.00 24.11	8
MOTA	878	N GLN A 112	44.101	31.156	58.772	1.00 24.94	7
MOTA	879		44.895	32.100	59.554		
ATOM	880					1.00 25.12	6
			45.082	33.413	58.779	1.00 25.14	6
ATOM	881	CG GLN A 112	45.545	33.224	57.330	1.00 28.51	6
MOTA	882	CD GLN A 112	45.789	34.534	56.594	1.00 29.13	6
ATOM	883	oel gln a 112	46.779	35.219	56.837	1.00 31.22	8
ATOM	884	NE2 GLN A 112	44.877	34.890	55.694	1.00 29.31	7
ATOM	885	C GLN A 112	44.107	32.362			
ATOM	886	O GLN A 112			60.827	1.00.24.62	6
			44.647	32.311	61.939	1.00 21.10	8
ATOM	887	N ALA A 113	42.813	32.622	60.644	1.00 24.41	7
ATOM-	888	CA ALA A 113	41.914	32.904	61.751	1.00 23.33	6
ATOM	889	CB ALA A 113	40.516	33.183	61.224	1.00 19.80	6
ATOM	890	C ALA A 113	41.901	31.733	62.729	1.00 25.34	
ATOM	891	O ALA A 113	41.925				6
	892			31.930	63.946	1.00 27.52	8
ATOM		N ILE A 114	41.859	30.509	62.211	1.00 24.39	7
ATOM	893	CA ILE A 114	41.867	29.356	63.106	1.00 24.49	6
ATOM	894	CB ILE A 114	41.524	28.042	62.371	1.00 23.46	6
ATOM	895	CG2 ILE A 114	41.902	26.855	63.227	1.00 18.97	6
ATOM	896	CG1 ILE A 114	40.030	28.015	62.034	1.00 21.17	6
ATOM	897	CD1 ILE A 114	39.598	26.791			
ATOM	898	C ILE A 114			61.239	1.00 22.51	6
			43.230	29.227	63.757	1.00 24.32	6
ATOM	899	O ILE A 114	43.328	28.817	64.907	1.00 24.74	8
ATOM	900	N GLU A 1:15	44.280	29.580	63.019	1.00 26.58	7
MOTA	901	CA GLU A 115	45.638	29.518	63.551	1.00 25.89	6
ATOM	902	CB GLU A 115	46.639	29.992	62.508	1.00 22.63	6
ATOM	903	CG GLU A 115	46.554	29.264	61.192		
ATOM	904	CD GLU A 115	17.668			1.00 20.39	6
				29.670	60.244	1.00 21.39	6
ATOM	905	OE1 GLU A 115	47.848	30.887	60.016	1.00 19.60	8
MOTA	906	OE2 GLU A 115	18.362	28.769	59.722	1.00 22.53	8
MOTA	907	C GLU A 115	45.724	30.422	64.774	1.00 27.56	6
ATOM	908	O GLU A 115	46.173	30.006	65.837	1.00 25.98	8
ATOM	909	N GLU A 116	45.267			1.00 31.19	7
ATOM	910	CA GLU A 116	45.282				
				32.631	65.705	1.00 35.80	6
ATOM	911	CB GLU A 116	44.676	33.959	65.237	1.00 36.91	6
MOTA	912	CG GLU A 116	45.434	34.605 ·	64.069	1.00 41.14	6
ATOM	913	CD GLU A 116 .	46.872	34.982	64.420	1.00 43.09	6
MOTA	914	OE1 GLU A 116	47.072	35.886	65.267	1.00 43.42	8
ATOM	915	CE2 GLU A 116	47.802	34.369	63.849		
	916					1.00 41.76	8
ATOM		•	44.543	32.131	66.947	1.00 35.11	6
MOTA	917	O GLU A 116	45.054	32.228	68.061	1.00 37.26	8
ATOM	918	N PHE A 117	43.343	31.598	66.761	1.00 34.30	7
ATOM	919	CA PHE A 117	42.577	31.096	67.893	1.00 34.44	6
ATOM	920	CB PHE A 117	41.300	30.399	67.415	1.00 35.45	6
	921	CG PHE A 117	40.383				
ATOM				29.979	68.533	1.00 37.14	6
ATOM	922	CD1 PHE A 117	39.705	30.930	69.290	1.00 35.80	6
ATOM	923	CD2 PHE A 117	40.196	28.630	68.832	1.00 41.05	.6
MOTA	924	CE1 PHE A 117	38.853	30.549	70.323	1.00 38.08	·6
		•			-		

		•				
ATOM	925 CE2 PHE A 117	. 30 3				
ATOM	11 11,	22.2				1 6
MOTA		38.66	58 29.19	8 70.61	7 1.00 38.64	
	- 1112 X 11/	43.42	30.09	4 68.669		
MCTA	928 O PHE A 117	43.49		£ 60.00		6
MOTA	929 N LEU A 118			• • .		8
ATOM	250 A 110	44.06		4 67.933	3 1.00 33.14	7
		44.89	8 28.15	8 68.523	3 1.00 32.62	
MCTA	45 PPO V 110	45.15				6
MOTA	932 CG LEU A 118	43.90				' 6
MOTA						6
ATOM	422 220 W 110	44.24	4 25.23	2 65.996	1.00 20.81	č
	934 CD2 LEU A 118	43.25	9 25.663	2 68.257		
MOTA	935 C LEU A 118	46.21				6
ATQM	936 O LEU A 118					6
MOTA	110	46.98			3 1.00 36.15	8
		46.48	1 29.974	68.843	1.00 34.75	_
ATOM	938 CA LYS A 119	47.67				
MOTA	939 CB LYS A 119	48.14				6
ATOM	940 CG LYS A 119					6
ATOM		48.61			1.00 37.98	6
		49.11.	1 32.430	66.263	1.00 43.40	
MOTA	942 CE LYS A 119	49.69			1.00 43.40	6
MOTA	943 NZ LYS A 119	50.16				6
ATOM	944 C LYS A 119				1.00 51.48	7
		47.27	31.191	70.705	1.00 34.85	. 6
ATOM	945 O LYS A 119	48.112	31.465		1 00 30 10	
ATOM	946 N GLY A 120	45.967				8
ATOM	947 CA GLY A 120					7
ATOM	948 C GLY A 120	45.431			1.00 36.25	6
		44.860	33.310	71.851	1.00 39.02	
ATOM	949 O GLY A 120	44.640	34.072	72.796		6
MOTA	950 N ASN A 121	44.619		70.750		8
ATOM	951 CA ASN A 121			70.586	1.00 38.48	7
ATOM	952 CB ASN A 121	44.079		70.247	1.00 37.47	6
ATOM		44.928	35.624	69.170	1.00 39.57	
	953 CG ASN A 121	46.340	35.871	69.622	1.00 41.81	6
ATOM	954 OD1 ASN A 121	47.078	34.938	69.926	1.00 41.81	5
ATOM	955 ND2 ASN A 121	46.727			1.00 47.67	8
ATOM	956 C ASN A 121			69.675	1.00 43.63	7
ATOM	957 O ASN A 121	42.637		69.772	1.00 36.59	6
		42.037	33.818	69.704	1.00 34.08	
ATOM	958 N VAL A 122	42.092	36.061	69.446	1.00 34.08	8
ATOM	959 CA VAL A 122	40.720	36.001		1.00 33.53	7
ATOM	960 CB VAL A 122		36.166	68.976	1.00 34.77	6
ATOM	961 CG1 VAL A 122	39.861	37.064	69.898	1.00 38.20	6
		38.418	37.096	69.388	1.00 37.55	
ATOM	962 CG2 VAL A 122	39.918	36.553	71.342	1 00 37.55	6
ATOM	963 C VAL A 122	40.731	36.781		1.00 37.77	6.
ATOM	964 O VAL A 122			67.596	1.00 31.08	6
ATOM		40.991	37. 967	67.441.	1.00 34.19	8
		40.451	35.975	66.588	1.00 31.14	7
ATOM	966 CA ALA A 123	40.451	36.476	65.231	1 00 30 06	
atom	967 CB ALA A 123	41.307	35.588		1.00 30.26	6
MCTA	968 C ALA A 123			64.327	1.00 32.14	6
ATOM		39.038	36.533	64.716	1.00 28.26	6
	·—··· ±25	38.132	35.924	65.281	1.00 29.28	8
ATOM	970 N PHE A 124	38.875	37.276	63.631	1.00 28.70	
atom	971 CA PHE A 124	37.601		63.031	1.00 28.70	7
MCTA	972 CB PHE A 124		37.475	62.976	1.00 28.38	6
ATOM	973 CG PHE A 124	36.920	38.713	63.563	1.00 29.16	6
		35.645	39.099	62.874	1.00 31.20	
ATOM	974 CD1 PHE A 124	34.679	38.139	62.564	7.00 31.20	6
MOTA	975 CD2 PHE A 124				1.00 32.00	6
MOTA	976 CE1 PHE A 124		40.435	62.579	1.00 29.53	6
TOM	977 CE2 PHE A 124	33.463	38.510	61.973	1.00 30.74	6
		34.165	40.813	61.988	1.00 27.45	
-TOM	.978 CZ PHE A 124	33.207	39.847	61.686	1 00 20 ==	6
TOM	979 C PHE A 124	37.880	77 (7:		1.00 28.72	6
NOT	980 O PHE A 124		37.671	61.496	1.00 30.19	6
		38.427	38.695	61.095	1.00 32.10	8
TOM	981 N ASN A 125	37.545	36.663	60.696	1.00 32.16	
TCM	982 CA ASN A 125	37.731	36.728	59.251	1 00 32.10	7
TOM	983 CB ASN A 125				1.00 30.10	6
TOM		38.247	35.393	58.712	1.00 32.02	6
		38.281	35.360	57.195	1.00 33.79	6
MCT.	985 OD1 ASN A 125	38.754	36.306	56.556		
MOT.	986 ND2 ASN A 125	37.790			1.00 31.85	8
TOM	987 C ASN A 125		34.268	56.609	1.00 30.51	7
TOM ·		36.403	37.083	58.584	1.00 29.80	6
	988 O ASN A 125	35.626	36.206	58.179	1.00 27.24	8
TOM	989 N PRO A 126	36.135	38.386	58.451		
TCM	990 CD PRO A 126	36.997	20.500	E0 022	1.00 28.25	7
		30.331	39.516	58.833	1.00 29.22	6

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MOTA	991	CA	PRO A 126	34.909	38.891	57.844	1.00 27.92	6
MOTA	992	CB	PRO A 126	35.139	40.407	57.856	1.00 29.07	6
MOTA	993	CG	PRO A 126	36.649	40.520	57.775	1.00 26.54	6
MOTA	994	С	PRO A 126	34.651	38.339	56.448	1.00 27.54	6
ATOM	995	0	PRO A 126	33.532	38.402	55.949	1.00 28.66	8
		N	ALA A 127	35.687	37.795	55.820	1.00 26.99	7
MOTA	996	14						
ATOM	997	CA	ALA A 127	35.548	37.244	54.477	1.00 26.54	6
	998	CB	ALA A 127	36.822	37.505	53.684	1.00 22.43	6
ATOM								
ATOM	999	С	ALA A·127	35.225	35.744	54.480	1.00 27.38	6
ATOM	1000	0	ALA A 127	35.038	35.140	53.423	1.00 29.04	8
MOTA	1001	N	GLY A 128	35.166	35.142	55.663	1.00 26.97	7
ATOM	1002	CA	GLY A 128	34.874	33.724	55.737	1.00 25.65	6
	1003	С	GLY A 128	33.389	33.486	55.880	1.00 26.17	6
MOTA								
ATOM.	1004	0	GLY A 128	32.600	34.428	55.804	1.00 27.39	8
MOTA	1005	N	GLY A 129	32.998	32.234	56.083	1.00 23.87	7
MOTA	1006	CA	GLY A 129	31.588	31.936	56.236	1.00 25.17	6
MOTA	1007	С	GLY A 129	30.847	31.674	54.937	1.00 25.88	6
MOTA	1008	0	GLY A 129	29:643	31.908	54.848	1.00 25.07	8
MOTA	1009	N	MET A 130	31.566	31.198	53.927	1.00 25.69	7
							•	
ATOM	1010	CA	MET A 130	30.981	30.872	52.622	1.00 26.48	6
ATOM	1011	CB	MET A 130	32.103	30.907	51.567	1.00 28.53	6
ATOM	1012	CG	MET A 130	32.795	32.288	51.467	1.00 26.54	б
		SD	MET A 130	34.413	32.366	50.613	1.00 26.29	16
ATOM	1013							
ATOM	1014	CE	MET A 130	34.080	31.512	49.062	1.00 25.85	6
ATOM	1015	С	MET A 130	30.355	29.463	52.768	1.00 24.47	6
	1016		MET A 130	30.761	28.502	52.113	1.00 17.67	8
ATOM		0						
MOTA	1017	N	HIS A 131	29.347	29.389	53.636	1.00 23.28	7
ATOM	1018	CA	HIS A 131	28.647	28.161	54.019	1.00 26.33	6
	1019	CB	HIS A 131	27.685	28.485	55.180	1.00 26.98	6
ATOM								
ATOM	1020	CG	HIS A 131	26.663	29.540	54.862	1.00 28.50	6
ATOM	1021	CD2	HIS A 131	26.225	30.030	53.677	1.00 28.65	б
MOTA	1022		HIS A 131	25.906	30.166	55.831	1.00 33.04	7
						55.259		
ATOM	1023		HIS A 131	25.051	30.995		1.00 27.75	6
ATOM	1024	NE2	HIS A 131	25.224	30. 9 32	53.952	1.00 26.97	7
ATOM	1025	С	HIS A 131	27.917	27.284	53.017	1.00 28.44	6
						53.390	1.00 31.15	٠ <u>a</u> ٠
MOTA	1026	0	HIS A 131	27.434	26.214			
ATOM	1027	И	HIS A 132	27.861	27.694	51.756	1.00 30.64	7
ATOM	1028	CA	HIS A 132	27.111	26.938	50.746	1.00 28.71	6
	1029		HIS A 132	26.321	27.941	49.890	1.00 27.21	6
MOTA		CB						
ATOM	1030	CG	HIS A 132	25.408	28.819	50.693	1.00 28.83	6
ATOM	1031	CD2	HIS A 132	25.111	30.137	50.578	1.00 28.92	6
	1032		HIS A 132	24.686	28.360	51.773	1.00 31.80	7
MOTA								
MOTA	1033		HIS A 132	23.981	29.353	52.285	1.00 29.95	6
ATOM	1034	NE2	HIS A 132	24.222	30.443	51.579	1.00 28.21	7
	1035	C	HIS A 132	27.889	25.970	49.851	1.00 28.51	6
ATOM								
MOTA	1036	0	HIS A 132	27.399.		49.533	1.00 23.44	8
MOTA	1037	N	ALA A 133	29.093	26.379	49.455	1.00 27.94	7
ATOM	1038	CA	ALA A 133	29.958	25.386	48.579	1.00 26.99	6
			ALA A 133	31.295	26.303	48.392	1.00 21.87	6
MOTA	1039	CB						
ATOM	1040	C	ALA A 133	30.199	24.164	49.078	1.00 26.69	6
MOTA	1041	0	ALA A 133	30.703	23.973	50.182	1.00 28.25	8
	1042		PHE A 134	29.850	23.174	48.255	1.00 26.73	7
ATOM		N						
ATOM	1043	CA	PHE A 134	30.046	21.773	48.615	1.00 25.04	6
MOTA	1044	CB	PHE A 134	29.070	20.855	47.875	1.00 19.20	6
	1045	CG	PHE A 134	27.629	21.199	48.100	1.00 15.75	6
ATOM						47.169	1.00 14.83	6
MOTA	1046	CDI	PHE A 134	26.929	21.960			2
MOTA	1047	CD2	PHE A 134	26.985	20.814	49.273	1.00 14.03	6
	1048	CE1	PHE A 134	25.614	22.336	47.404	1.00 14.84	6
MOTA		C = 2	DUE 3 134	25.670		49.519	1.00 12.07	6
MOTA	1049		PHE A 134		21.184			
ATOM	1050	CZ	PHE A 134	24.985	21.949	48.581	1.00 14.59	6.
ATOM	1051	С	PHE A 134	31.460	21.310	48.319	1.00 29.48	6
			PHE A 134	32.291	22.066	47.822	1.00 33.19	8
ATOM	1052	0						7
ATOM	1053	N	LYS A 135	31.713	20.045	48.620	1.00 31.52	
ATOM	1054	CA	LYS A 135	33.012	19.427	48.427	1.00 29.15	6
	1055	CB	LYS A 135	32.923	17.971	48.885	1.00 29.45	6
ATCM			LYS A 135		17.131	48.638	1.00 32.46	6
ATOM	1056	CG	PID W IDD	34.152	エト・アコエ	20.030	JZ. 140	-

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MOTA	105	7 C	D LYS A 135		33.965	15.734	49,221	1 00 20 67	_
ATOM	105				34.234		50.716		6
ATOM	105	9 N	Z LYS A 135		35.679		51.001	1.00 26.25	6 7
MOTA	1060	0 C	LYS A 135		33.513		46.993	1.00 30.22	6
ATOM	106		LYS A 135		34.714		46.763	1.00 30.00	8
ATOM	1062				32.600	19.434	46.028	1.00 31.69	7
ATOM	1063	-			32.995	19.489	44.619	1.00 32.88	6
ATOM	1064				33.038	18.077	44.040	1.00 31.41	6
ATOM	1065				33.882	17.241	44.810	1.00 35.41	8
ATOM	1066 1067				32.097	20.347	43.727	1.00 33.55	6
ATOM ATOM	1068		SER A 136 ARG A 137		31.921	20.031	42.553	1.00 36.11	8
ATOM	1069				31.536	21.425	44.262	1.00 30.61	7
ATOM					30.664 29.324	22.272 21.554	43.459	1.00 32.28	6
ATOM	1071				28.224	22.458	43.202 42.627	1.00 35.91	6
ATOM	1072				26.819	21.836	42.751	1.00 43.90 1.00 48.28	6
ATOM	1073				26.571	20.767	41.787	1.00 48.28	6
ATOM	1074	CZ			26.150	20.960	40.538	1.00 55.30	7 6
ATOM	1075		11 ARG A 137		25.921	22.185	40.090	1.00 54.06	7
MOTA	1076		12 ARG A 137		25.969	19.922	39.728	1.00 58.96	7
MOTA	1077		ARG A 137		30.405	23.631	44.113	1.00 30.24	6
ATOM	1078		ARG A 137		30.380	23.748	45.338	1.00 23.11	8
MOTA	1079 1080		ALA A 138		30.219	24.653	43.279.		7
ATOM ATOM	1080				29.944	26.000	43.757	1.00 27.36	6
MOTA	1082		ALA A 138		30.149 28.496	26.997	42.645	1.00 27.57	6
ATOM	1083		ALA A 138		27.747	26.003 25.083	44.213 43.865	1.00 26.45	6
ATOM	1084	N	ASN A 139		28.090	27.021	44.975	1.00 27.30 1.00 22.47	8
ATOM	1085	CA			26.711	27.063	45.471	1.00 23.85	7 6
ATOM	1086	CB	ASN A 139		26.406	25.738	46.218	1.00 16.82	6
MOTA	1087	CG			25.040	25.718	46.900	1.00 14.45	6
ATOM	1088		1 ASN A 139		24.019	26.084	46.319	1.00 13.39	8
ATOM	1089	ND			25.018	25.249	48.139	1.00 20.08	7
ATOM ATOM	1090 1091	C	ASN A 139		26.444	28.277	46.368	1.00 26.09	6
MOTA	1091	N	ASN A 139 GLY A 140		27.239	28.600	47.260	1.00 27.50	8
ATOM	1093	CA	GLY A 140		25.326 24.965	28.954 30.106	46.114	1.00 24.83	7
ATOM	1094	c	GLY A 140		25.991	31.211	46.916 46.890	1.00 22.24 1.00 22.35	6
MOTA	1095	0	GLY A 140		26.256		47.910	1.00 22.35	6 8
ATOM	1096	N	PHE A 141		26.570	31.437	45.717	1.00 25.60	7
MOTA	1097	CA	PHE A 141		27.582	32.476	45.518	1.00 26.47	6
ATOM	1098	CB	PHE A 141		27.204	33.765	46.258	1.00 28.05	6
ATOM	1099	CG	PHE A 141		25.925	34.391	45.792	1.00 28.61	6
ATOM	1100 1101	CD2	PHE A 141		25.352	35.428	46.518	1.00 30.74	6
MOTA MO A	1101	CEI			25.312	33.975	44.620	1.00 29.10	6
A-COM	1103	CE2		د	24.193 24.150	36.044	46.087	1.00 29.33	6
ATCM	1104	CZ	PHE A 141	-	23.589	34.583 35.621	44.177 44.912	1.00 31.03 1.00 32.59	6
MOTA	1105	·C	PHE A 141		28.954	32.038	45.991	1.00 32.59	6 6
MOTA	1106	0	PHE A 141		29.938	32.727	45:733	1.00 29.72	8
MOTA	1107	N	CYS A 142		29.025		46.667	1.00 21.11	7
MOTA	1108	CA	CYS A 142		30.296	30.399	47.192	1.00 22.30	6
MOTA	1109	CB	CYS A 142		30.062	29.787	48.567	1.00 21.31	6
ATOM	1110	SG	CYS A 142		28.943	30.748	49.582	1.00 22.93	16
MOTA	1111 1112	c	CYS A 142		31.017	29.366	46.326	1.00 22.13	6
MOTA MOTA	1113	O N	CYS A 142		30.408	28.389	45.878	1.00 22.97	8
ATOM	1114	CA	TYR A 143 TYR A 143		32.317 33.129	29.573 28.632	46.111	1.00 23.09	7
ATOM	1115	CB	TYR A 143		34.063	29.365	45.335 44.375	1.00 23.05 1.00 21.60	6 6
ATOM	1116	CG	TYR A 143		33.377	30.379	44.373	1.00 21.60	6
ATOM	1117		TYR A 143		32.969	31.609	43.999	1.00 23.29	6
ATOM	1118		TYR A 143		32.365	32.555	43.199	1.00 23.26	6
MOTA	1119		TYR A 143		33.154	30.117	42.135	1.00 22.52	6
ATOM	1120		TYR A 143		32.544	31.061	41.317	1.00 24.82	6
ATOM	1121	CZ	TYR A 143		32.153	32.281	41.857	1.00 27.55	6
ATOM	1122	OH	TYR A 143		31.553	33.241	41.064	1.00 32.35	8

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i

	1123	С	TYR A 14	1	33.960	27.766	46.290	1.00 24.22	6
MOTA					34.266	26.606	45.998	1.00 24.58	8
ATOM	1124	0	TYR A 14				47.437	1.00 23.83	7
MOTA	1125	N	ILE A 14	i	34.327	28.329			
ATOM	1126	CA	ILE A 14	l	35.086	27.566	48.425	1.00 20.24	6
ATOM	1127	CB	ILE A 14		36.547	27.982	48.453	1.00 17.27	6
	1128	CG2	ILE A 14		37.231	27.354	49.662	1.00 11.03	6
MOTA					37.185	27.603	47.110	1.00 14.93	6
MOTA	1129	_	ILE A 14				46.946	1.00 19.68	6
ATOM	1130		ILE A 14		38.601	28.028			
MOTA	1131	С	ILE A 14	l	34.495	27.703	49.815	1.00 21.77	6
ATOM	1132	0	ILE A 14	l .	34.288	28.811	50.318	1.00 21.19	8
MOTA	1133	N	ASN A 14		34.212	26.555	50.424	1.00 23.00	7
	1134	CA	ASN A 14		33.616	26.508	51.750	1.00 20.92	6.
ATOM			ASN A 14		32.902	25.170	51.935	1.00 17.08	6
MOTA	1135	CB			32.079	25.125	53.203	1.00 21.04	6
MOTA	1136	CG	ASN A 14						
MOTA	1137		ASN A 14		32.549	25.508	54.276	1.00 20.97	8
ATOM	1138	ND2	ASN A 14	5	30.844	24.640	53.093	1.00 20.93	7
ATOM	1139	С	ASN A 14	5	34.706	26.669	52.806	1.00 19.68	6
ATOM	1140	0	ASN A 14	5	35.201	25.679	53.351	1.00 20.64	. 8
	1141	N	ASN A 14		35.079	27.911	53.100	1.00 16.28	7
ATOM			ASN A 14		36.123	28.143	54.088	1.00 19.34	6
MOTA	1142	CA			36.428	29.651	54.207	1.00 20.27	6
ATOM	1143	CB	ASN A 14						6
MOTA	1144	CG	ASN A 14		35.292	30.444	54.795	1.00 18.05	
MOTA	1145	OD1	ASN A 14	5	35.079	30.421	55.999	1.00 25.83	8
ATOM	1146	ND2	ASN A 14	S	34.552	31.149	53.948	1.00 16.04	7
ATOM	1147	С	ASN A 14	5 .	35.775	27.504	55.443	1.00 20.48	6
	1148	0	ASN A 14		36.663	27.027	56.151	1.00 19.88	8
ATOM	1149	N	PRO A 14		34.482	27.485	55.819	1.00 19.38	7
ATOM			PRO A 14		33.312	28.068	55.135	1.00 17.48	6
ATOM	1150	CD			34.058	26.877	57.087	1.00 22.25	6
MOTA	1151	CA	PRO A 14					1.00 20.15	6
ATOM	1152	CB	PRO A 14		32.539	27.065	57.057		
ATOM	1153	CG	PRO A 14	7	32.407	28.378	56.305	1.00 20.81	6
ATOM	1154	С	PRO A 14	7	34.443	25.383	57.188	1.00 26.89	6
ATOM	1155	0	PRO A 14		35.066	24.954.	58.169	1.00 29.10	8
	1156	N	ALA A 14		34.070	24.596	56.176	1.00 25.88	7
ATOM	1157	CA	ALA A 14		34.372	23.164	56.174	1.00 25.47	6
MOTA			ALA A 14		33.670	22.468	55.009	1.00 21.84	6
ATOM	1158	CB			35.870	22.916	56.100	1.00 25.94	6
ATOM	1159	C	ALA A 14					1.00 27.19	8
ATOM	1160	0	ALA A 14		36.382	21.971	56.701		7
MOTA	1161	N	VAL A 14		36.574	23.756	55.349	1.00 26.11	
ATOM	1162	CA	VAL A 14	9	38.017	23.609	55.233	1.00 24.04	` 6
ATOM	1163	CB	VAL A 14	9	38.622	24.663	54.267	1.00 26.16	6
ATOM	1164	CG1	VAL A 14	9	40.135	24.476	54.158	1.00 25.36	6
	1165		VAL A 14	9	37.970	24.544	52.886	1.00 26.81	6
ATOM	1166	C	VAL A 14		38.516	23.870	56.640	1.00 23.57	6
MOTA					39.453	23.228	57.122	1.00 19.75	8
MOTA	1167	0	VAL A 14			24.815	57.299	1.00 22.20	7
MOTA	1168	N	GLY A 15		37.850			1.00 25.43	6
ATOM	1169	CA	GLY A 15		38.210	25.175	58.654		
ATOM	1170	C	GLY A 15	0	38.130	23.975	59.568	1.00 27.19	6
ATOM	1171	0	GLY A 15	0	39.112	23.620	60.221	1.00 27.05	8
ATOM	1172	N	ILE A 15	1	36.959	23.348	59.618	1.00 25.56	7
	1173	CA	ILE A 15		36.775	22.176	60.457	1.00 28.24	6
ATOM			ILE A 15		35.317	21.654	60.389	.1.00 29.41	6
ATOM	1174	CB			35.251	20.215	60.869	1.00 26.02	6
ATOM	1175	CG2	ILE A 15					1.00 33.31	6
MOTA	1176	CG1	ILE A 15	Ţ	34.394	22.540			
MOTA	1177	CD1	ILE A 15	1	34.255	23.967	60.759	1.00 36.83	6
ATOM	1178	С	ILE A 15	1	37.723	21.039	60.075	1.00 29.32	6
	1179	Ō	ILE A 15		38.340	20.420	60.947	1.00 29.08	8
MOTA	1180	N	GLU A 15			20.769	58.778	1.00 29.91	7
MOTA			GLU A 15		38.704	19.690	58.315	1.00 32.58	6
ATOM	1181	CA			38.575	19.526	56.802	1.00 35.07	б
ATCM	1182	CB	GLU A 15				56.393	1.00 38.51	6
ATOM	1183	CG	GLU A 15		37.269	18.848			6
MOTA	1184	CD	GLU A 15		37.120		56.987	1.00 41.41	
ATOM	1185	OE1	GLU A 15		36.089		56.718	1.00 45.64	8
	1186	OE2	GLU A 15		38.030	16.992	57.723	1.00 40.09	8
ATOM	1187	C	GLU A 15	2	40.145		58.721	1.00 32.97	6
ATOM	1188	0	GLU A 15		40.879		58.978	1.00 30.15	8
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Figure 17-19

ATOM	1189	N TYR A 153	40.5	1 21.170) E0 766		
ATOM	1190	CA TYR A 153	41.8				-
MOTA	1191	CB TYR A 153	42.03				
ATOM	1192	CG TYR A 153	43.28				6
ATOM	1193	CD1 TYR A 153	44.49				
ATOM	1194	CE1 TYR A 153	45.65				6
ATOM	1195	CD2 TYR A 153	43.25				6
MCTA		CE2 TYR A 153	44.38				
ATOM		CZ TYR A 153	45.58				
ATOM	1198	OH .TYR A 153	46.69				6
ATOM		TYR A 153	41.91				8
ATOM	1200 (TYR A 153	42.86				6
ATOM	1201	V LEU A 154	40.86				8
ATOM		CA LEU A 154	40.73				7
MOTA		B LEU A 154	39.44		62.823		6
ATOM	1204	G LEU A 154	39.39	9 23.407			6
ATOM	1205	D1 LEU A 154	37.99				6.
ATOM	1206	D2 LEU A 154	40.41				6
ATOM	1207 0		40.73			1.00 24.95	6
MCTA	1208 0		41.22			1.00 29.56	6
ATOM	1209 N		40.17			1.00 28.36	8
ATOM	1210 C	A ARG A 155	40.13		62.499	1.00 31.95	7
ATOM	1211 C	B ARG A 155	39.12		61.561	1.00 33.00	6
MOTA	1212 C	G ARG A 155	37.70		61.769	1.00 33.13	6
ATOM	1213 C	D ARG A 155	. 36.67		60.863	1.00 32.84	6
MOTA	1214 N	E ARG A 155	36.15		61.363	1.00 32.92 1.00 33.98	6
ATOM	1215 C	Z ARG A 155	35.19		60.741	1.00 33.98	7
ATOM	1216 N	H1 ARG A 155	34.67		59.605	1.00 37.33	6
ATOM		H2 ARG A 155	34.732		61.259	1.00 38.67	7 7
ATOM	1218 C		41.52		62.331	1.00 33.97	6
ATOM	1219 o		41.869		62.985	1.00 32.95	8
MOTA	1220 N		42.318	3 17.548	61.467	1.00 34.20	7
ATOM	1221 C		43.679		61.243	1.00 36.32	6
ATOM	1222 CI		44.249		59.942	1.00 37.57	6
ATOM	1223 CC		45.673	17.187	59.638	1.00 40.32	6
MOTA	1224 CI		46.116	17.532	58.220	1.00 40.33	6
MOTA	1225 CI		45.180		57.184	1.00 41.27	6
ATOM	1226 NZ 1227 C		45.015		57.364	1.00 37.92	7
MCTA MCTA	1227 C 1228 O	LYS A 156	44.539		62.428	1.00 36.17	6
MCTA	1229 N	LYS A 156	45.582		62.699	1.00 34.53	. 8
MOTA	1230 CA	LYS A 157 LYS A 157	44.093		63.132	1.00 36.71	7
MOTA	1231 CE		44.820		64.294	1.00 37.09	6
MOTA	1232 CG		44.495		64.566	1.00 37.02	6
MOTA	1233 CD		44.982		63.477	1.00 36.22	6
MCTA	1234 TE		46.468		63.239	1.00 37.91	6
MOTA	1235 JZ		46.993 48.434	22.100	62.107	1.00 39.35	6
MCTA	1236	LYS A 157		21.815	61.842	1.00 38.78	7
ATOM	1237 0	LYS A 157	44.498 45.204	18.178	65.515	1.00 35.61	6
ATCM	1238 N	GLY A 158	43.433	18.232	66.518	1.00 36.38	8
MOTA	1239 CA		43.097	17.392	65.431	1.00 34.37	7
ATCM	1240 C	GLY A 158	41.782	16.537 16.781	66.552	1.00 38.08	6
ATCM	1241 0	GLY A 158	41.460	16.053	67.267	1.00 38.78	6
ATOM	1242 N	PHE A 159	41.023	17.791	68.208 66.855	1.00 41.07	8
MOTA	1243 CA	PHE A 159	39.743	18.046		1.00 36.75	7
ATOM	1244 CB	PHE A 159	39.246	19.459	67.505 67.213	1.00 33.83	6
MCTA	1245 CG	PHE A 159	40.115	20.521	67.787	1.00 32.65	6
ATOM		l PHE A 159	41.404	20.724	67.297	1.00 29.97	6
MOTA		PHE A 159	39.672	21.289	68.853	1.00 30.20 1.00 29.28	6
MCTA		PHE A 159	42.241	21.289	67.862	1.00 29.28	6
MOTA	1249 CE2		40.498	22.246	69.428	1.00 28.96	6
ATCM	1250 CZ	PHE A 159	41.785	22.442	68.931	1.00 29.67	6 6
ATOM	1251 C	PHE A 159	38.732	17.026	67.025	1.00 33.41	6
MCTA	1252 0	PHE A 159	38.664	16.716	65.838	1.00 31.61	8
ATCM	1253 N	LYS A 160	37.951	16.506	67.966	1.00 35.13	7
MOTA	1254 CA	LYS A 160	36:947	15.493	67.677	1.00 35.39	6
		• .	_		//	22.33	

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ATOM	1255	CE	LYS A 160		37.342	14.198	68.389	1 00	36.43	•
MOTA	1256	CG	LYS A 160		38.535		67.708			6
ATOM	1257	CD			39.312			1.00	40.67	6
ATOM	1258	CE					68.615	1.00	44.68	6
					38.425		69.345	1.00	49.23	6
ATOM	1259	NZ			37.593		70.411		50.63	7
MOTA	1260	С	LYS A 160		35.524	15.927	68.027	1.00	35.94	6
MOTA	1261	0	LYS A 160		34.561	15.241	67.691		35.72	8
ATOM	1262	N	ARG A 161	•	35.399		68.718		34.35	7
ATOM	1263	CA	ARG A 161		34.091	17.618	69.044		34.95	
ATOM	1264	C3	ARG A 161		33.771			1.00	34.95	6
ATOM	1265	CG	ARG A 161			17.525	70.535	1.00	33.94	6
	1266			•	33.427	16.132	70.992	1.00	38.25	6
MOTA		CD	ARG A 161		32.823	16.131	72.386	1.00	41.17	б
ATOM	1267	NE	ARG A 161		33.719	16.722	73.378	1.00	47.64	7
MOTA	1268	cz	ARG A 161		34.912	16.233	73.705		47.92	6
MOTA	1269	NH:	l ARG A 161		35.372	15.131	73.121		47.56	7
MOTA	1270	NH2	2 ARG A 161		35.648	16.858	74.516	1 00	46.95	
ATOM	1271	С	ARG A 161		34.113	19.076	68.598	1.00	40.55	7
ATOM	1272	ō	ARG A 161		34.468			1.00	34.58	6
	1273	N				19.980	69.357		33.77	8
MOTA			ILE A 162		33.741	19.280	67.341	1.00	31.74	7
MOTA	1274	CA	ILE A 162		33.735	20.594	66.735	1.00	29.83	· 6
ATOM	1275	CB	ILE A 162		34.429	20.542	65.362		29.96	6
ATOM	1276	CG2	! ILE A 162		34.580	21.942	64.784		30.57	6
ATOM	1277	CG1	ILE A 162		35.801	19.891	65.522		28.81	6
ATOM	1278	CDI	ILE A 162		36.537	19.685	64.224		33.05	
ATOM	1279	C	ILE A 162		32.300	21.050				6
ATOM	1280	ŏ	ILE A 162				66.560		29.66	6
	1281	N	LEU A 163		31.416	20.241	66.266	1.00	25.24	8
ATOM					32.081	22.351	66.745		30.00	7
MOTA	1282	CA	LEU A 163	•	30.754	22.945	66.617	1.00	30.48	6
ATOM	1283	CB	LEU A 163		30.236	23.406	67.992	1.00	32.25	6
MOTA	1284	CG	LEU A 163		28.934	24.229	68.044	1.00	31.21	6
ATOM	1285	CD1	LEU · A 163		27.804	23.494	67.326		31.58	6
ATOM	1286	CD2	LEU A 163		28.569	24.502	69.493		25.00	6
ATOM	1287	С	LEU A 163		30.717	24.122	65.659		29.23	
ATOM	1288	0	LEU A 163		31.596	24.980	65.654			6
ATOM		N	TYR A 164		29.675				29.72	8
ATOM		CA	TYR A 164			24.157	64.846		29.68	7
		CB			29.500	25.244	63.899		29.89	6
ATOM			TYR A 164		29.512	24.688	62.470		27.81	6
MOTA		CG	TYR A 164		29.377	25.742	61.399	1.00	27.79	6
MOTA		CD1	TYR A 164		30.390	26.670	61.168	1.00	24.82	6
MOTA		CE1	TYR A 164		30.247	27.655	60.198	1.00	24.51	6
MOTA		CD2	TYR A 164		28.216	25.827	60.631	1.00	27.61	6
ATOM	1296	CE2	TYR A 164		28.065	26.808	59.662		25.67	5
ATOM	1297	CZ	TYR A 164		29.078	27.718	59.451		25.63	6
MOTA	1298	OH	TYR A 164		28.898	28.704	58.506	1 00	27.10	
ATOM		c	TYR A 164		28.149	25.907	64.218		28.38	8
ATOM		ō	TYR A 164		27.119					6
ATOM		N	ILE A 165			25.225	64.277		29.43	8
		CA			28.166	27.217	64.464		24.30	7
ATOM			ILE A 165		26.941		64.754	1.00	22.93	6
MOTA		CB	ILE A 165		26.985	28.649	66.143	1.00	22.00	6
ATOM		CG2	ILE A 165		25.765	29.559	66.312	1.00	16.15	6
ATOM		CG1	ILE A 165		27.033	27.567	67.240	1.00	20.78	
ATOM	1306	CD1	ILE A. 165		27.185	28.101	68.650	1.00	15.49	6
MOTA		2	ILE A 165		26.784	29.010	63.657		24.45	6
MOTA)	ILE A 165		27.605	29.921·	63.506		23.17	
ATOM		V	ASP A 166		25.709					8
ATOM			ASP A 166			28.871	62.895		24.20	7
					25.478	29.726	61.749		20.78	6
ATOM			ASP A 166		25.314	28.809	60.548		17.64	6
ATOM			ASP A 166		25.410	29.529	59.256	1.00	19.93	6
atom			ASP A 166		24.536	30.391	59.004		20.20	8
MOTA			ASP A 166		26.366	29.231	58.491		17.64	8
ATOM	1315		ASP A 166		24.290	30,670	61.895		22.79	6
ATOM	1316		ASP A 166		23.134	30.256	61.826	1.00	22.70	. 8
ATOM	1317 N		LEU A 167		24.583					
			LEU A 167			31.952	62.085		25.40	7
ATOM ·					23.536	32.954	62.250	1.00	25.76	6
MOTA			LEU A 167		23.963	33.991	63.288		26.65	6
MOTA	1320, C	. ي	LEU A 167		24.364	33.463	64.674	1.00	26.75	6

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MOTA	1321	CD	1 LEU A 167		24.741	34.647	65.552	1.00 26.24	6
ATOM	1322		2 LEU A 167		23.225				
						32.661	65.302	1.00 23.45	6
MOTA	1323	С	LEU A 167		23.162	33.660	60.951	1.00 26.37	6
MOTA	1324	0	LEU A 167		22.386	34.613	60.971	1.00 25.95	8
MOTA	1325	N	ASP A 168		23.726	33.208	59.828	1.00 29.66	Ž.
ATOM-	1326	CA	ASP A 168		23.410	33.787			
							58.520	1.00 28.35	6
MOTA	1327	CB	ASP A 168		24.057	32.987	57.390	1.00 33.29	6
MOTA	1328	CG	ASP A 168		23.937	33.676	56.037	1.00 35.38	6
MOTA	1329	OD	L ASP A -168		24.892	34.388	55.659	1.00 39.48	8
MOTA	1330		2-ASP A 168		22.893	33.531	55.364		
								1.00 33.40	8
ATOM	1331	С	ASP A 168		21.906	33.614	58.408	1.00 28.74	6
ATOM	1332	0	ASP A 168		21.354	32.648	58.948	1.00 26.21	8
ATOM	1333	N	ALA A 169	-	21.239	34.524	57.711-	1.00 26.16	7
ATOM	1334	CA	ALA A 169		19.793	34.415	57.579	1.00 24.39	6
ATOM	1335	CB	ALA A 169		19.233	35.640	56.879	1.00 22.75	
									6
MOTA	1336	C	ALA A 169		19.420	33.157	56.813	1.00 24.37	6 -
ATOM	1337	0	ALA A 169		18.266	32.752	56.824	1.00 22.34	8
ATOM	1338	N	HIS A 170		20.405	32.542	56.156	1.00 25.78	7
MOTA	1339	CA	HIS A 170		20.180	31.327	55.375	1.00 25.20	6
ATOM	1340	CB	HIS A 170		20.667	31.501	53.936	1.00 25.76	6
ATOM	1341	CG	HIS A 170		20.122				
						32.711	53.245	1.00 29.08	6
MOTA	1342		HIS A 170		19.338	32.834	52.147	1.00 30.59	6
ATOM	1343	NDI	. HIS A 170		20.384	33.995	53.675	1.00 30.77	7
MOTA	1344	CE1	. HIS A 170		19.784	34.858	52.873	1.00 29.07	6
ATOM	1345	NE2	HIS A 170		19.143	34.180	51.939	1.00 32.19	7
ATOM	1346	C	HIS A 170		20.895	30.113	55.958	1.00 26.00	6
ATOM	1347	ō	HIS A 170		21.913	30.234			
	•						56.637	1.00 25.76	8
ATOM	1348	N	HIS A 171		20.349	28.939	55.658	1.00 27.29	7
MOTA	1349	CA	HIS A 171		20.893	27.655	56.090	1.00 25.01	6
MOTA	1350	CB	HIS A 171		19.934	26.532	55.663	1.00 24.93	6
MOTA	1351	CG	HIS A 171		20.468	25.148	55.889	1.00 26.56	6
MOTA	1352	CD2	HIS A 171		20.674	24.123	55.028	1.00 22.34	6
ATOM	1353		HIS A 171		20.823	24.678	57.137	1.00 25.35	7
ATOM	1354		HIS A 171		21.222	23.424		1.00 22.68	6
ATOM	1355		HIS A 171		21.140				
						23.062	55.767	1.00 24.13	7
MOTA	1356	C	HIS A 171		22.267	27.413	55.471	1.00 24.74	6
MOTA	1357	0	HIS A 171		22.540	27.863	54.356	1.00 28.22	8
ATOM	1358	N	CYS A 172		23.131	26.705	56.190	1.00 23.03	7
ATOM	1359	CA	CYS A 172		24.467	26.389	55.683	1.00 23.41	6
ATOM	1360	CB	CYS A 172		25.497	26.474	56.812	1.00 19.31	6
ATOM	1361	SG	CYS A 172		25.005	25.631	58.318	1.00 16.78	16
ATOM	1362	С	CYS A 172		24.484	24.997	55.048	1.00 25.45	6
	1363	õ	CYS A 172						
MOTA					25.203	24.098	55.483	1.00 24.47	8
MOTA	1364	N	ASP A 173		23.664	24.839	54.015	1.00 26.67	7
ATOM	1365	CA	ASP A 173		23.542	23.593	53.269	1.00 26.47	6
MOTA	1366	CB	ASP A 173		22.735	23.857	51.993	1.00 26.33	6
ATOM	1367	CG	ASP A 11.3		23.281	25.030	51.179	1.00 27.06	6
ATOM	1368	OD1	'ASP A 173	-	22.539	25.558	50.330	1.00 23.43	8
ATOM	1369		ASP A 173		24.454	25.417	51.372	1.00 29.38	8
ATOM	1370	C	ASP A 173						
					24.872	22.932	52.922	1.00 26.65	6
ATOM	1371	0	ASP A 173		24.940	21.708	52.784	1.00 28.38	8
ATOM	1372	N	GLY A 174		25.926	23.737	52.793	1.00 25.24	7
ATOM	1373	CA	GLY A 174		27.227	23.198	52.447	1.00 23.11	6
MOTA	1374	C	GLY A 174		27.896	22.505	53.612	1.00 25.64	6
ATOM	1375	Õ	GLY A 174		28.443	21.408	53.462	1.00 27.67	8
	1376	N	VAL A 175		27.848	23.144	54.778		7
ATCM								1.00 24.29	
ATOM	1377	CA	VAL A 175		28.459	22.602	55.989	1.00 22.20	6
ATOM	1378	CB	VAL A 175		28.536	23.672	57.101	1.00 20.15	6
ATOM	1379	CG1	VAL A 175		29.449	23.192	58.218	1.00 20.11	6
MOTA	1380		VAL A 175		29.015	24.989	56.530	1.00 18.74	6
ATOM	1381	С	VAL A 175		27.647	21.409	56.505	1.00 22.85	6
	1382	0	VAL A 175		28.173	20.512	57.173	1.00 20.07	8
ATOM			GLN A 176		26.356		_		7
ATOM	1383	N				21.404	56.203	1.00 24.12	
atom ·	1384	CA	GLN A 176		25.518	20.303	56.629	1.00 27.18	6
MOTA	1385	CB	GLN A 176		24.045	20.611	56.355	1.00 32.86	6
MOTA	1386	CG	GLN A 176		23.084	19.483	56.726	1.00 36.04	6
			•				•		

ATOM	138'	7 C	D GLN A 176		21.62	2.0	19.86			
ATOM	1388	8 0	E1 GLN A 176		21.11					36 6
ATOM	1389	9 N	E2 GLN A 176		20.93		20.782			59 8
ATOM	1390) c			25.95		19.15			
ATOM	1391	L O			26.32		19.083			70 6
ATOM	1392						18.066			39 8
ATOM					25.95		19.194			96 7
ATOM					26.34		18.062			L6 6
MOTA					26.39		18.460			37 6
ATOM					26.35		17.256		1.00 36.2	20 6
ATOM			E1 GLU A 177		26.27		17.626		1.00 40.7	70 6
ATOM			E2 GLU A 177		27.32		17.967		1.00 46.7	78 8
ATOM			GLU A 177		25.15		17.590		1.00 39.3	9 8
ATOM					27.70		17.516	54.137	1.00 31.6	6
ATOM	1401		GLU A 177		27.86		16.317		1.00 32.8	11 8
ATOM	1402	C.	ALA A 178		28.66		18.419		1.00 33.3	9 7
ATOM	1403				30.02		18.072	54.673	1.00 31.6	3 6
ATOM	1404	CE			30.83		19.338	54.856	1.00 30.9	6 6
ATOM		C	ALA A 178		30.20		17.185		1.00 30.6	
	1405	0	ALA A 178		31.03	2	16.276	55.876	1.00 27.9	5 B
ATOM	1406	N	PHE A 179		29.44		17.444	56.961	1.00 31.0	1 7
ATOM	1407	CA			29.59	0	16.656	58.184	1.00 31.3	4 6
MOTA	1408	CB			30.14	7	17.532	59.310	1.00 30.1	
ATOM	1409	CG			31.189	9	18.505	58.858	1.00 27.7	
ATOM	1410	CD			30.827	7	19.790	58.466	1.00 28.2	
ATOM	1411	CD	2 PHE A 179		32.522		18.124	58.766	1.00 28.3	4 6
ATOM	1412		1 PHE A 179		31.778	В	20.688	57.988	1.00 26.6	3 6
ATOM	1413	CE			33.487		19.013	58.285	1.00 28.7	
ATOM	1414	CZ	PHE A 179		33.111		20.300	57.895	1.00 28.7	9 6
ATOM	1415	С	PHE A 179		28.300		16.003	58.664	1.00 28.6	
MOTA	1416	0	PHE A 179		28.218		15.542	59.803	1.00 32.0	
MOTA	1417	N	TYR A 180		27.305		15.960		1.00 30.5	
ATOM	1418	CA	TYR A 180		26.001		15.377	57.787	1.00 34.2	5 7
MOTA	1419	CB	TYR A 180		25.062			58.099	1.00 38.6	
ATOM	1420	CG	TYR A 180 .		23.593		15.605	56.911	1.00 38.9	
ATOM	1421	CDI					15:453	57.220	1.00 37.9	16
ATOM	1422	CEI			22.938		14.232	57.064	1.00 35.83	36
ATOM	1423	CD2			21.589		14.103	57.373	1.00 39.20	0 6
ATOM	1424	CE2	TYR A 180		22.861		16.543	57.694	1.00 37.5	
ATOM	1425	CZ	TYR A 180		21.518		16.430	58.007	1.00 40.28	3 6
ATOM	1426	ОН	TYR A 180		20.882		15.211	57.848	1.00 41.92	2 6
ATOM	1427	C	TYR A 180		19.549		15.110	58.188	1.00 43.43	1 8
ATOM	1428	ō	TYR A 180		26.133		13.884	58.382	1.00 40.28	3 6
MOTA	1429	N	ASP A 181		25.158		13.192	58.680	1.00 39.27	7 8
ATOM	1430	CA	ASP A 181		27.363		13.402	58.319	1.00 43.51	
ATOM	1431	CB		-	27.638		11.994	58.519	1.00 45.89	
ATOM	1432	CG	ASP A 181		28.414		11.487	57.303	1.00 51.00	6.
ATOM	1433	OD1	ASP A 181		28.830		10.050	57.436	1.00 56.84	6
ATOM	1434				29.637		9.750	58.345	1.00 59.47	
	1435		ASP A 181		28.348		9.221	56.629	1.00 60.73	
ATOM		C	ASP A 181		8.398		11.665	59.804	1.00 44.75	6
ATOM	1436	0	ASP A 181		8.257	1	10.568	60.350	1.00 44.69	8
ATOM		N	THR A 182	2	9.194	1	12.506	60.298	1.00 41.26	7
MOTA		CA	THR A 182	2	9.975	1	12.337	61.495	1.00 39.51	
ATOM		CB	THR A 182	3	1.408		12.881	61.355	1.00 39.19	
MOTA	1440	CG1	THR A 182	3	2.171		2.508	62.505	1.00 37.82	
ATOM	1441	CG2	THR A 182		1.395		4.397	61.232	1.00 40.12	8
ATOM		C	THR A 182		9.370	1	2.910	62.759		
atom	1443	0	THR A 182		8.609		3.876	62.716	1.00 38.58	. 6
ATOM	1444	N	ASP A 183		9.712		2.304	63.890	1.00 41.24	
ATOM	1445	CA	ASP A 183		9.211				1.00 37.39	7
ATOM			ASP A 183		8.824		2.773	65.171	1.00 39.24	
MOTA			ASP A 183		0.010		1.588	66.061	1.00 40.31	6
ATOM		OD1	ASP A 183				0.723	66.433	1.00 41.64	6
ATOM	1449	2סכ	ASP A 183		0.725		0.268	65.520	1.00 42.53	8
ATOM		2	ASP A 183		0.221		0.494	67.640	1.00 42.46	8
ATOM		5	ASP A 183		0.286		3.621	65.853	1.00 40.34	6
ATCM					0.109		4.071	66.983	1.00 42.07	8
-1 011	1734 I	٠.	GLN A 184	3	1.400	1	3.830	65.154	1.00 39.29	7

MOTA	1453	CA	GLN A 184		32.50	6 14.63	35 65.67	1 1 00 2- 0-	
ATOM	1454	CB	GLN A 184		33.83				6
MOTA	1455	CG	GLN A 184		34.22				' 6
ATOM	1456		GLN A 184		35.59				
MOTA	1457	OE1	GLN A 184		35.85				
MOTA	1458	NE2			36.49				8
ATOM	1459	С	GLN A 184		32.22				
MOTA	1460	0	GLN A 184		32.80	3 16.99			6
MOTA	1461	N	VAL A 185		31.32			/.121	. 8
MOTA	1462	CA	VAL A 185		30.98				
ATOM	1463	CB	VAL A 185		31.30				
ATOM	1464	CG1	VAL A 185		31.00				6
ATOM	1465	CG2	VAL A 185		32.77				6
MOTA	1466	C	VAL A 185		29.50	17.97			6
ATOM	1467	0	VAL A 185		28.68				6
MOTA	1468	N	PHE A 186		29.18				8
ATOM	1469	CA	PHE A 186		27.798			7. 1.00 31.44	7
ATOM	1470	CB	PHE A 186		27.524			1.00 30.96	.6
MOTA	1471	CG	PHE A 186	•	26.059				6
MOTA	1472		PHE A 186		25.552			1.00 30.54	6
ATOM	1473	CD2	PHE A 186		25.179				6 6
MOTA	1474	CE1	PHE A 186		24.191	18.95		1.00 33.28	6
MOTA	1475	CE2			23.815	20.45			6
ATOM	1476	CZ	PHE A 186		23.318	19.73			6
ATOM	1477	С	PHE A 186		27.490	20.798		1.00 30.37	6
MOTA	1478	0	PHE A 186		28.189		64.751	1.00 31.32	8
ATOM ATOM	1479 1480	И	VAL A 187		26.435			1.00 31 14	7
MOTA	1481	CA CB	VAL A 187		26.024			1.00 32.05	6
ATOM	1482		VAL A 187 VAL A 187		26.018			1.00 33.54	6
ATOM	1483		VAL A 187		25.574			1.00 32.07	ε
ATOM	1484	C	VAL A 187		27.420			1.00 35.44	6
ATOM	1485	ō	VAL A 187		24.638	22.439			6
ATOM	1486	N	LEU A 188		23.666	21.686			8
ATOM	1487		LEU A 188		24.579 23.336	23.638		1.00 29.44	7
ATOM	1488	CB	LEU A 188		23.433	24.228		1.00 29.39	6
MOTA	1489		LEU A 188		22.293	24.665 25.589		1.00 29.52	6
MOTA	1490		LEU A 188		20.970	24.844		1.00 27.92	5
ATOM	1491	CD2	LEU A 188			26.107		1.00 25.87	6
ATOM	1492	C	LEU A 188		23.161	25.454		1.00 27.69 1.00 31.89	6
ATOM	1493	0	LEU A 188		24.130	26.175		1.00 31.50	6
MOTA	1494		SER A 189		21.929	25.700	63.250	1.00 29.93	8 7
ATOM	1495		SER A 189		21.682	26.831	62.390	1.00 24.65	6
ATOM	1496	CB :	SER A 189		21.873	26.411	60.942	1.00 22.40	6
ATOM	1497		SER A 189		21.585	27.485	60.083	1.00 19.12	8
ATOM	1498		SER A 189		20.516	27.462	62.540	1.00 27.00	6
ATOM	1499		SER A 189		1996	26.774	62.577	1.00 26.72	8
ATOM	1500 1501		LEU A 190		20.521	28.783	62.669	1.00 27.41	7
atom Atom			LEU A 190		19.096	29.554	62.735	1.00 29.68	6
MOTA		_	EU A 190		19.185	30.682	63.771	1.00 29.84	6
ATOM			EU A 190 EU A 190		19.108	30.366		1.00 26.79	6
ATOM			EU A 190		19.020	31.662	66.045	1.00 23.44	6
ATOM			EU A 190 EU A 190		17.881	29.549	65.546	1.00 27.63	6
ATOM			EU A 190		19.046	30.141	61.329	1.00 29.58	6
MOTA			IS A 191		20.084 17.864	30.525		1.00 32.40	8
ATOM			IS A 191		17.766	30.206	60.727	1.00 29.61	7
MOTA			IS A 191		18.595	30.726	59.368	1.00 29.72	6
MOTA			IS A 191		18.225	29.839	58.432	1.00 26.47	6
MOTA			IS A 191		18.918	28.392	58.504	1.00 28.18	6
ATOM			IS A 191		16.989	27.313	58.940	1.00 28.88	6
ATCM			IS A 191		16.938	27.921	58.118	1.00 31.00	7
			IS A 191		18.095	26.614 26.220	58.312	1.00 30.54	6
			IS A 191		16.329	30.812	58.810 58.856	1.00 27.21	7
			IS A 191			30.411	59.535	1.00 28.05	6
	1518 N		LN A 192		16.183	31.346	57.649	1.00 27.81	8
	-					JI. J40	37.049	1.00 29.39	7

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  ATOM
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                                                                  1.00 22.22
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                                                                 1.00 23.38
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          1528
                 ÇA
                     SER A 193
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                                                                       31.52
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                                                                      37.57
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                                                                 1.00
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                                                        53.403
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                CG
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                CA
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                        A 195
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                                                       53.527
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               CE1
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                                     11.637
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                                                       54.706
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                                                                     33.10
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                   TYR A 196
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                                                       54.776
                                                                1.00 36.75
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 ATOM
               CE2
                    TYR A 196
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                                              31.825
                                                       55.968
                                                                1.00
                                                                     35.26
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               CZ
                    TYR A 196
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11.704
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               ОН
 ATOM
                    TYR A 196
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                                                       57.087.
                                                                1.00 37.09
                                                                               8
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                    TYR A 196
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                                                       51.188
                                                                1.00 34.89
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        1560
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                   TYR A 196
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                                     14.297
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        1569
              CG
                   PHE A 198
                                    19.810
                                                      48.578
                                             30.013
                                                                1.00 43.59
                                                                               б
              CD1
ATOM
        1570
                   PHE A 198
                                                      47.325
                                    19.783
                                             29.404
                                                               1.00 44.74
                                                                               6
                   PHE A 198
ATOM
        1571
              CD2
                                    20.970
                                             29.929
                                                      49.336
                                                               1.00 41.86
                                                                              6
ATOM
        1572
              CE1
                   PHE A 198
                                    20.894
                                             28.729
                                                      46.833
                                                               1.00 41.42
                                                                              6
ATOM
        1573
              CE2
                   PHE A 198
                                    22.079
                                                      48.849
                                             29..251
                                                               1.00 43.30
                                                                              6
MOTA
        1574
              CZ
                   PHE A 198
                                    22.040
                                                      47.595
                                             28.652
                                                               1.00 41.86
                                                                              6
ATOM
       1575
              С
                   PHE A 198
                                    18.139
                                             29.140
                                                      50.967
                                                               1.00
                                                                     37.00
                                                                              6
       1576
              0
                                             29.754
ATOM
                   PHE A 198
                                    18.166
                                                      52.036
                                                               1.00
                                                                     36.43
                                                                              8
       1577
              N
                   PRO A 199
MOTA
                                    18.641
                                                      50.848
                                                               1.00 37.63
                                             27.892
       1578
              CD
                   PRO A 199
ATOM
                                    19.298
                                                      51.997
                                             27.238
                                                               1.00
                                                                     35.29
                                                                              6
       1579
              CA
                   PRO A 199
ATOM
                                    18.727
                                             27.008
                                                      49.673
                                                               1.00
                                                                     36.52
                                                                              6
       1580
              CB
                   PRO A 199
                                    19.702
MOTA
                                                      50.138
                                             25.936
                                                               1.00 34.96
       1581
              CG
                  PRO A 199
ATOM
                                    19.281
                                                      51.565
                                             25.770
                                                               1.00 34.57
                                                                              б
              С
                  PRO A 199
ATOM
       1582
                                    17.409
                                                      49.222
                                                               1.00 35.72
                                             26.380
                                                                              б
ATOM
       1583
              0
                  PRO A 199
                                    17.386
                                                      48.225
                                                               1.00 37.36
                                            25.663
                                                                              8
       1584
             N
                  PHE A 200
ATOM
                                    16.331
                                                      49.962
                                             26.638
                                                               1.00 33.78
```

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Figure 17-25

		•
ATO	M 1585 CA PHE A 200	15.004 26.090 49.662 1.00.32 15 6
ATO		49.002 1.00 32.15
ATO		10:222 1.00 28.39 6
		14.600 27.827 47.025 1.00.00 00
ATO		15.749 78 385 47 386 1 66 64 65
ATO		13 466 28 623 47 866 1 00 20
ATO:		15 767 29 712 46 992 1 00 25 04 6
ATO:		25.712 40.882 1.00 25.68 6
ATO		1.00 27.03 6
ATO	M 1593 C PHE A 200	14.626 30.498 47.013 1.00 24.90 6
-	200	14.947 24.574 49.842 1.00 32.66 6
ATO		13.925 24.033 50.264 1 00 31 22 0
ATO		16 0/17 22 000 40 400 4 55 5
ATOI	M 1596 CA GLU A 201	15 120 22 420 40 505
ATO	M 1597 CB GLU A 201	1 1 1 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
ATOI	1 1598 CG GLU A 201	1,00 32.30
ATON		16.879 22.182 47.175 1.00 33.52 6
		18.012 21.864 46.232 1.00 34.56 6
ATO1		18.396 20.678 46.117 1 00 36 35 8
ATOP		18 523 22 814 45 605 1 60 35
ATOM	1 1602 C GLU A 201	16 760 21 011 50 000
ATOM	1 1603 O GLU A 201	15 535
ATOM		100 20.31
ATOM		17.511 22.239 51.566 1.00 31.64 7
ATOM	502	17.795 21.780 52.917 1.00 32.34 6
	202	19.276 21.432 53.092 1.00 36.91 6
ATOM		10 700 20 226 52 205
ATOM	202	20 212 20 500 50 001
ATOM	1609 CE LYS A 202	20 052 10 400 50 000
ATOM	1610 NZ LYS A 202	21 504 20 20 20 20 20 20 20 20 20 20 20 20 20
ATOM		
ATOM		17.421 22.849 53.937 1.00 30.55 6
ATOM		· ±0.8// 23.890 53.586 1 00 26 55 0
	021 11 205	17.710 22.571 55.203 1.00 30.59 7
ATOM	1614 CA GLY A 203	17.422 23.519 56.259 1.00 30.24 6
ATOM	1615 C GLY A 203	16 216 22 210 57 125
MOTA	1616 O GLY A 203	16 016 02 000 000 000
ATOM	1617 N PHE A 204	15 526 20 101
ATOM	1618 CA PHE A 204	14 744 01 000 50 100
ATOM	1619 CB PHE A 204	14.344 21.779 57.657 1.00 23.25 6
ATOM	1620 CG PHE A 204	13.366 20.917 56.863 1.00 21.25 6
ATOM	1621 CD1 PHE A 204	12.855 21.573 55.635 1.00 18.60 6
		13.605 21.560 54.461 1.00 16.43 6
ATOM	1622 CD2 PHE A 204	11.654 22.273 55.664 1.00 14.82 6
ATOM	1623 CE1 PHE A 204	13 160 22 245 52 222 4 22
MOTA	1624 CE2 PHE A 204	11 206 22 262 24 244
ATOM	1625 CZ PHE A 204	11 005
ATOM	1626 C PHE A 204	14 626 24 251
ATOM	1627 O PHE A 204	20.575 1.00 25.72
ATOM	1628 N LEU A 205	15.578 20.318 59.118 1.00 22.68 8
ATOM		13.760 21.376 59.942 1.00 20.94 7
		13.877 20.818 61.272 1.00 24.83 6
MOTA	1630 CB LEU A 205	12.678 21.259 52.110 1.00 21.29 6
MOTA	1631 CG LEU A 205	12 672 20 011 : 2 550 1 22 22
ATOM	1632 CD1 LEU A 205	14 011 24 100
ATOM	1633 CD2 LEU A 205	11 479 21 456 64 255
ATOM	1634 C LEU A 205	14 002 10 202 61 202
ATOM	1635 O LEU A 205	
ATOM	1636 N GLU A 206	14.443 18.730 62.310 1.00 28.59 8
		13.625 18.628 60.211 1.00 33.52 7
ATOM	1637 CA GLU A 206	13.693 17.166 60.142 1.00 39.79 6
ATOM	1638 CB GLU A 206	12 776 16 666 50 000
ATOM	1639 CG GLU A 206	11 001
ATOM	1640 CD GLU A 206	11 014 10 000
ATOM	1641 OE1 GLU A 206	0.000
ATOM	1642 OE2 GLU A 206	11 070
ATOM		11.839 18.786 57.661 1.00 56.48 8
		13.114 16.674 59.847 1.00 40.00 6
ATOM	1644 O GLU A 206	15.483 15.541 60.180 1.00 39.35 8
ATOM ·	1645 N GLU A 207	15.903 17.536 59.217 1.00 39.38 7
MOTA	1646 CA GLU A 207	45 000
ATOM	1647 CB GLU A 207	12 226
ATOM	1648 CG GLU A 207	16 000
ATOM	1649 CD GLU A 207	16 070
		16.978 19.452 55.773 1.00 38.16 6
MOTA	1650 OE1 GLU A 207	18.071 20.016 55.537 1.00 35.44 8

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	MOTA MOTA MOTA	165: 165: 165:	2 C 3 O	GLU	J A 207 J A 207 J A 207	15.870 18.139 18.560	17.239 18.303	60.134 60.590	1.00	34.62 36.18 34.81	. 8 6 8
	MOTA MOTA	1654 1655			A 208	18.381				34.45	7
	ATOM	1656			A 208 A 208	19.164 18.260				38.53	6
	ATOM	1657	7 C	32 ILE	A 208	19.097				41.12 41.36	6 6
	MOTA	1658			A 208	17.193				42.21	6
	MOTA	1659			A .208	16.291				44.81	6
	MOTA MOTA	1660 1661			A 208	20.407				36.66	6
	ATOM	1662			A 208 A 209	21.243 20.540				34.03	8
	ATOM	1663			A 209	21.703				36.80	7 6
	ATOM.	1664			A 209	21.509		59.805		40.93	6
	ATOM	1665			A 209	20.477		59.145	1.00	40.26	8
	ATOM ATOM	1666 1667			A 210 A 210	22.508		59.775		42.16	7
	ATOM	1668			A 210	22.492 22.810		58.930 57.488		43.30	6
	ATOM	1669			A 210	22.826		56.478		47.08	6 6
	MOTA	1670			A 210	23.256	9.915	55.089		56.27	6
	ATOM	1671	OE	1 GLU	A 210	24.412	10.371	54.941	1.00	56.19	8
	MOTA MOTA	1672 1673			A 210 A 210	22.437	9.826	54.145		60.28	8
	ATOM	1674			A 210	23.583 24.750	9.276 9.457	59.473 59.152		41.47 43.97	6
	ATOM	1675		GLY	A 211	23.203	8.307	60.299		39.97	8 7
	ATOM	1676			A 211	24.181	7.405	60.885		37.34	6
	ATOM	1677 1678	C		A 211	24.642	7.952	62.224		37.84	6
	ATOM ATOM	1679	O N		A 211 A 212	23.820 25.948	8.408 7.910	63.019		37.30	8
	ATOM	1680	CA		A 212	26.490	8.440	62.485 63.733		38.52 38.29	7 6
	ATOM	1681	CB	LYS	A 212	28.020	8.359	63.731		40.54	6
	ATOM	1682	CG		A 212	28.570	6.950	63.675		46.39	6
	MOTA MOTA	1683 1684	CD		A 212 A 212	28.149 28.556	6.147	64.910		51.59	6
	ATOM	1685	NZ		A 212	30.030	4.676 4.478	64.809 64.662		52.77 55.48	6 7
	MOTA	1686	С	LYS	A 212	26.061	9.897	63.866		37.68	6
	ATOM	1687	0		A 212	25.814	10.389	64.962		34.75	8
	ATOM ATOM	1688 1689	N CA		A 213 A 213	25.956 25.577	10.574	62.728		38.89	7
	MOTA	1690	c		A 213	24.126	11.975 12.295	62.724 63.020		43.58 43.99	6 6
	ATOM	1691	0	GLY	A 213	23.737	13.464	63.024		44.67	8
	ATOM	1692	N		A 214	23.321	11.268	63.265	1.00	46.02	7
	MOTA MOTA	1693 1694	CA CB		A 214 A 214	21.907	11.467	63.562		45.61	6
	MOTA	1695	CG		A 214	21.168 19.675	10.130 10.252	63.469 63.249		47.77 49.25	6 6
	MOTA	1696	CD	LYS .	A 214	19.078	8.901	62.911		51.64	6
	· MOTA	1697	CE		A 214	17.637	9.038	62.440	1.00	54.30	6
	MOTA MOTA	1698 1699	NZ C		A 214 A 214	17.030	7.727	62.034		56.09	7
	ATOM	1700	ō		A 214	21.809 22.210	12.047 11.410	64.970 65.942		44.22 45.04	6 8
	ATOM	1701	N		A 215	21.292	13.266			42.89	7
	MOTA	1702	CA	GLY A		21.193	13.904	66.373		40.20	6
	ATOM	1703 1704	С	GLY A		22.295	14.931	66.600		39.90	6
	MOTA MOTA	1704	N N	GLY A		22.356 23.175	15.548	67.668		40.12	8
	ATOM	1706	CA	TYR A		24.261	15.111 16.086	65.612 65.726	1.00	38.49 35 91	7 6
	ATOM	1707	CB	TYR A	216	25.632	15.421	65.618	1.00		6
	MOTA	1708	CG	TYR A		25.935	14.461	66.738	1.00	39.54	6
	MOTA	1709 1710		TYR A		25.296	13.220	66.810	1.00		6
-	ATOM ATOM	1711		TYR A		25.561 26.852	12.333	67.849 67.739	1.00		6 6
	ATOM	1712	CE2	TYR A	216	27.124	14.795 13.915	68.786	1.00		6
	ATOM	1713	CZ	TYR A	216	26.475	12.685	68.833	1.00	41.86	6
	ATOM	1714	OH	TYR A		26.743	11.802	69.852	1.00	43.04	8
	ATOM	1715 1716	С 0	TYR A		24.182	17.215	64.709	1.00		6
	ATOM	1,10	٠.	IIV W	. 410	25.194	17.832	64.375	1.00	33.37	8
			€								

ATO	M 1717 N ASN A 217	00.00				
ATO		22.97		– —		, ,
ATO		22.72		63.26		6
		22.69				- 6
ATO:		22.45	7 19.177	60.82		E
ATO		21.35	4 19.705	60.719	9 1.00 25.00	8
ATO:		23.50	1 19.558			7
ATO		21.36			1.00 29.09	
ATO	M 1724 O ASN A 217	20.43			5 1.00 26.93	6
ATO	M 1725 N LEU A 218	21.26			1.00 26.93	8
ATO		20.01				7
ATO		20.02				6
ATO		18.72				6
· ATO						6
ATON		19.100				6
ATON		17.87				6
ATON	1 1732 O LEU A 218	19.789			1.00 25.04	6
		20.596		63.415	1.00 25.23	8
ATOM		18.681	. 22.436			7
ATOM		18.310	23.636	61.829		6
ATOM		17.809	23.298	60.417	1.00 25.69	
ATOM		18.748		59.646	1.00 26.10	6
ATOM		19.927		59.505	1.00 28.53	6
ATOM		18.220		59.114		8
ATOM		17.129	24.248	62.582		7
ATOM		16.373				6
ATOM		16.952		63.246	1.00 34.84	8
ATOM				62.472		7
ATOM		15.826		63.129	1.00 32.50	6
ATOM		16.259		64.350	1.00 32.32	6
ATOM		15.029		65.014	1.00 29.46	6
ATOM		16.978	26.160	65.374	1.00 29.65	6
	1747 C ILE A 220	16.080	25.138	66.027	1.00 28.65	6
MOTA		15.140	27.106	62.123	1.00 35.36	6
ATOM	1748 O ILE A 220	15.469	28.290	62.009	1.00 35.52	8
ATOM	1749 N PRO A 221	14.185	26.553	61.359	1.00 36.87	7
ATOM	1750 CD PRO A 221 .	13.718	25.158	61.359	1.00 35.12	6
ATOM	1751 CA PRO A 221	13.445	27.318	60.356	1.00 35.41	6
ATOM	1752 CB PRO A 221	12.509	26.262	59.767	1.00 35.68	
ATOM	1753 CG PRO A 221	13.319	24.992	59.911	1.00 33.86	6 6
MOTA	1754 C PRO A 221	12.696	28.437	61.053	1.00 34.37	
MOTA	1755 O PRO A 221	12.014	28.199	62.043		6
ATOM	1756 N LEU A 222	12.815	29.655		1.00 38.79	8
ATOM	1757 CA LEU A 222	12.138	30.796	60.547	1.00 34.76	7
ATOM	1758 CB LEU A 222	13.173	31.735	61.166	1.00 33.87	6
ATOM	1759 CG LEU A 222	14.104		61.798	1.00 35.13	6
ATOM	1760 CD1 LEU A 222	15.234	31.163	62.876	1.00 33.07	6
ATOM	1761 CD2 LEU A 222		32.150	63.154	1.00 34.04	6
ATOM	1762 C LEU A 222	13.312	30.856	64.141	1.00 32.39	6
ATOM	1763 O LEU A 222	11.287	31.567	60.157	1.00 32.15	6
ATOM	1764 N PRO A 223	11.669	31.740	59.000	1. 0 31.32	- 8
		10.127	32.060	60.601	1.17 30.97	7
MOTA		9.606	31.913	61.972	1.00 32.34	6
ATOM		9.173	32.818	59.789	1.00 30.55	6
MOTA	1767 CB PRO A 223	7.957	32.893	60.702	1.00 29.44	6
ATOM	1768 CG PRO A 223	8.626	33.068	62.046	1.00 31.02	6
ATOM	1769 C PRO A 223	9.645	34.205	59.366	1.00 29.20	6
ATOM	1770 O PRO A 223	10.694	34.680	59.796	1.00 31.95	8
MOTA	1771 N LYS A 224	8.841	34.841	58.521	1.00 26.14	7
ATOM	1772 CA LYS A 224	9.115		58.026	1.00 23.54	
MOTA	1773 CB LYS A 224	8.285	36.443	56.766	1.00 24:71	6
ATOM	1774 CG LYS A 224	8.563		55.619	1.00 23.83	6
ATOM	1775 CD LYS A 224	7.737		54.394	1 00 22.83	6
ATOM	1776 CE LYS A 224	8.065			1.00 20.59	6
ATOM	1777 NZ LYS A 224	7.198		53.329	1.00 27.03	6
ATOM	1778 C LYS A 224			52.122	1.00 30.48	7
ATOM	1779 O LYS A 224			59.111	1.00 25.48	6
		7.999		60.055	1.00 22.67	8
ATOM				58.960	1.00 26.88	7
ATOM	1781 CA GLY A 225			59.925	1.00 29.80	6
ATCM	1782 C GLY A 225	9.396	39.188 (51.286	1.00 32.96	6
•			-			

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MOTA	1783	O GLY A	225	9.068	39.861	62.271	1.00 31.20	8
ATOM	1784			10.299	38.216	61.338	1.00 32.86	7
MOTA	1785			10.235	37.877	62.575	1.00 34.55	6
ATOM	1786			12.149	36.958	62.255	1.00 34.46	6
MOTA	1787			12.143	36.413	63.407	1.00 34.48	6
ATOM	1788			12.146	35.425	64.212		
ATOM	1789						1.00 33.18	6
ATOM	1790			14.207	35.724	62.847	1.00 31.39	6
ATOM	1791	C LEU A O LEU A		11.481	39.160	63.255	1.00 36.29	6
	1792			12.156	39.970	62.613	1.00 33.87	8
MOTA			205	11.131	39.358	64.531	1.00 37.31	7
MOTA	1793	CA ASN A		11.592	40.536	65.279	1.00 37.26	6
MOTA	1794	CB ASN A		10.444	41.212	66.053	1.00 35.57	6
MOTA	1795	CG ASN A		9.920	40.368	67.208	1.00 36.07	6
ATOM	1796	OD1 ASN A		10.678	39.940	68.089	1.00 35.08	8
ATOM	1797	ND2 ASN A		8.611	40.143	67.218	1.00 32.33	7
ATOM	1798	C ASN A		12.688	40.096	66.250	1.00 37.95	6
ATOM	1799	O ASN A		12.869	38.890	66.473	1.00 37.08	.8
ATOM	1800	N ASP A		13.403	41.063	66.832	1.00 36.07	7
ATOM	1801	CA ASP A		14.505	40.754	67.751	1.00 37.63	6
ATOM	1802	CB ASP A		14.996	42.007	68.486	1.00 36.48	
MOTA	1803	CG ASP A		15.480	43.088	67.545	1.00 37.52	6
MOTA	1804	OD1 ASP A		15.936	42.752	66.427	1.00 35.28	8
ATOM	1805	OD2 ASP A		15.426	44.274	67.937	1.00 39.01	8
ATOM	1806	C ASP A		14.204	39.678	68.783	1.00 37.56	6
ATOM	1807	O ASP A		14.921	38.678	68.869	1.00 39.53	8
ATOM	1808	N ASN A		13.155	39.889	69.572	1.00 38.37	7
ATOM	1809	CA ASN A		12.766	38.935	70.605	1.00 37.49	6
MOTA	1810	CB ASN A		11.422	39.352	71.200	1.00 37.38	6
ATOM	1811	CG ASN A		11.490	40.709	71.877	1.00 40.47	6
ATOM	1812	OD1 ASN A		12.041	40.840	72.973	1.00 41.76	8
MOTA	1813	ND2 ASN A		10.960	41.735	71.212	1.00 36.50	7
ATOM	1814	C ASN A		12.680	37.530	70.017	1.00 37.64	6
ATOM	1815	O ASN A		13.446	36.634	70.395	1.00 35.76	8
ATOM	1816	N GLU A		11.758	37.351	69.076	1.00 36.01	7
MOTA	1817	CA GLU A		11.574	36.062	68.425	1.00 34.74	6
ATOM	1818	CB GLU A		10.753	36.242	67.153	1.00 35.55	6
ATOM	1819			9.382	36.820	67.407	1.00 36.95	6
ATOM	1820 1821	CD GLU A OE1 GLU A		8.580	36.960	66.144	1.00 35.30	6
ATOM	1822	OE2 GLU A		9.042	37.670	65.229	1.00 36.98	8
ATOM	1823	C GLU A		7.490 12.916	36.361	66.065	1.00 36.71	8
ATOM	1824	O GLU A		13.143	35.421	68.082	1.00 33.92	6
ATOM ATOM	1825	N PHE A		13.143	34.238	68.346	1.00 32.74	8
ATOM	1826	CA PHE A		15.116	36.207 35.712	67.487 67.123	1.00 32.03 1.00 30.55	7 6
ATOM	1827	CB PHE A		15.932	36.821	66.460	1.00 33.86	6
ATOM	1828	CG 'HE A		17.295	36.381	66.012	1.00 35.86	. 6
ATOM	1829	CD1 -HE A		17.438	35.334	65.102	1.00 40.41	6
ATOM	1830	CD2 HE A		18.436	37.021	66.480	1.00 36.58	6
ATOM	1831	CE1 PHE A		18.709	34.932	64.661	1.00 43.00	6
ATOM	1832	CE2 PHE A		19.711	36.632	66.049	1.00 39.07	6
ATOM	1833	CZ PHE A		19.849	35.586		1.00 40.52	6
ATOM	1834	C PHE A		15.835	35.232		1.00 30.63	6
ATOM	1835	O PHE A		16.177	34.042	68.497	1.00 29.66	8
ATOM	1836	N LEU A		16.049	36.162	69.310	1.00 24.94	7
	1837	CA LEU A		16.742	35.857	70.556	1.00 22.82	6
ATOM	1838	CB LEU A		16.724	37.084		1.00 24.96	. 6
MOTA	1839	CG LEU A		17.507		71.468		
MOTA	1840	CD1 LEU A		17.316	38.282	70.890	1.00 29.34	6
ATOM		CD2 LEU A		18.991	39.549	71.746	1.00 24.38	6
MOTA	1841			16.150	37.903	70.787	1.00 27.39	6
ATOM	1842	C LEU A			34.638	71.261	1.00 22.44	6
MOTA	1843			16.882	33.793	71.767	1.00 20.37	8 7
MOTA	1844	N PHE A		14.825 14.131	34.552	71.289	1.00 24.81	
MOTA	1845			12.623	33.422	71.905	1.00 25.81	6 6
MOTA	1846				33.535	71.641	1.00 24.37	6
ATOM	1847	CG PHE A		11.811	32.373	72.157	1.00 24.18	6
ATOM	1848	CD1 PHE A	درے	11.491	32.264	73.503	1.00 25.59	O

ATOM 1849 CD2 PHE A 233	11 320 31 200 0
ATOM 1850 CE1 PHE A 233	11.339 31.397 71.284 1.00 25.75 6 10.698 31.198 73.974 1.00 25.23
ATOM 1851 CE2 PHE A 233	10 7.00 25.33 6
ATOM 1852 CZ PHE A 233	10 000 11.00. 24.02 6
ATOM 1853 C PHE A 233	14 000 11.00 44.77 6
ATOM- 1854 O PHE A 233	10 00 -
ATOM 1855 N ALA A 234	14 504 8
ATOM 1856 CA ALA A 234	15 202
ATOM 1857 CB ALA A 234	14 727
ATOM 1858 C ALA A 234	15 75 5 5 5 5 5 5 6 5 6 6 6 6 6 6 6 6 6 6
ATOM 1859 O ALA A 234	10.303 30.645 69.433 1.00 30.76 6
ATOM 1860 N LEU A 235	12 262 272 05.400 1.00 30.04 8
ATOM 1861 CA LEU A 235	10
ATOM 1862 CB LEU A 235	10 540
ATOM 1863 CG LEU A 235	19.348 32.819 69.703 1.00 34.12 6
ATOM 1864 CD1 LEU A 235	21.039 32.745 69.316 1.00 36.33 6
ATOM 1865 CD2 LEU A 235	21.025 34.156 69.205 1.00 36.44 6
ATOM 1866 C LEU A 235	21.803 31.939 70.330 1.00 35.64 6
ATOM 1867 O LEU A 235	18.970 30.846 71.176 1.00 30.75 6
ATOM 1868 N GLU A 236	19.048 29.835 71.312 1.00 30.12 8
ATOM 1869 CA GLU A 236	10.347 31.435 72.192 1.00 29.03 7
ATOM 1870 CB GLU A 236	10.418 30.931 73.561 1.00 33.32 6
ATOM 1871 CG GLU A 236	17.479 31.730 74.452 1.00 35.06 6
ATOM 1872 CD GLU A 236	17.043 33.176 74.635 1.00 42.35 6
ATOM 1873 OE1 GLU A 236	15.610 34.022 74.843 1.00 47.12 6
ATOM 1874 OE2 GLU A 236	15.686 33.557 75.556 1.00 48.91 8
ATOM 1875 C GLU A 236	10.372 35.150 74.297 1.00 49.07 8
ATOM 1876 O GLU A 236	17.366 29.473 73.639 1.00 34.65 6
ATOM 1877 N LYS A 237	16.715 28.593 74.116 1.00 30.43 8
ATOM 1878 CA LYS A 237	16.767 29.250 73.176 1.00 35.67 7
ATOM 1879 CB LYS A 237	16.138 27.943 73.175 1.00 35.51 6
ATOM 1880 CG LYS A 237	14.791 28.060 72.452 1.00 37.01 6 13.745 27.032 72.848 1.00 37.05
ATOM 1881 CD LYS A 237	10 710 -1002 72.040 1.00 39.05 6
ATOM 1882 CE LYS A 237	17 212
ATOM 1883 NZ LYS A 237	10 000 0000
ATOM 1884 C LYS A 237	15 22
ATOM 1885 O LYS A 237	12.403 1.00 35.07 6
ATOM 1886 N SER A 238	17 = ,5.001 1.00 30.60 8
ATOM 1887 CA SER A 238	10 000 /
ATOM 1888 CB SER A 238	10 4-10 52.10
ATOM 1889 OG SER A 238	19 014 28 160 00 000 1.00 32.01 6
ATOM 1890 C SER A 238	
ATOM 1891 O SER A 238	20 21.34
ATOM 1892 N LEU A 239	20 20 20 20 20
ATOM 1893 CA LEU A 239	21 660 26 650 70 11.00 31.19
ATOM 1894 CB LEU A 239	20 20-
ATOM 1895 CG LEU A 239	20 20.03
ATOM 1896 CD1 LEU A 239	22 24 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2
ATOM 1897 CD2 LEU A 239	23 663 29 691 70 770
	25 1.00 23.11
ATOM 1899 O LEU A 239	'- '- '- '- '- '- '- '- '- '- '- '- '
ATOM 1900 N GLU A 240	20 367 25 025 74 025
ATOM 1901 CA GLU A 240	20 094 24 055 75 136 1 33.86 /
ATOM 1902 CB GLU A 240	18 799 25 260 75 040 4 00
ATOM 1903 CG GLU A 240	10 500 43.21 6
ATOM 1904 CD GLU A 240	19 677 24 703 70 000
ATOM 1905 OE1 GLU A 240	19 969 25 300 75 700
ATOM 1906 OE2 GLU A 240	20 21 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2
ATOM 1907 C GLU A 240	20 22 -0.504 70.055 1.00 55.12 8
ATOM 1908 O GLU A 240	20 532 22 22 25
ATOM 1909 N ILE A 241	19 423 23 23 23 2
ATOM 1910 CA ILE A 241	19.310 21.896 73.035 1.00 38.00
ATOM 1911 CB ILE A 241	10 465
ATCM · 1912 CG2 ILE A 241	18 536 30 505 31
ATOM 1913 CG1 ILE A 241	17 (1)2 22 224 22 22 2
ATOM 1914 CD1 ILE A 241	16.147 22.478 70.843 1.00 27.53 6
	•

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ATOM	1915	C	ILE A 241	20.713	21.372	72.747	1.00 39.56	6
ATOM	1916		ILE A 241	20.984		72.936		
	1917		VAL A 242					8
MOTA				21.605		72.299		7
ATOM	1918			22.979	21.842	72.015	1.00 45.09	6
ATOM	1919	CE	3 VAL A 242	23.808	22.959	71.329	1.00 45.76	6
ATOM	1920	CG	31 VAL A 242	25.242	22.479	71.116	1.00 43.09	6
ATOM	1921	CG	2 VAL A 242	23.182	23.334	69.991	1.00 46.41	6
ATOM	1922		VAL A 242	23.698	21.453	73.300	1.00 45.69	
	1923		VAL A 242					6
ATOM				24.191	20.331	73.423	1.00 46.30	8
MOTA	1924		LYS A 243	23.750	22.373	74.259	1.00 44.60	7
ATOM	1925	CA		24.427	22.088	75.513	1.00 46.96	6
ATOM	1926	CE	LYS A 243	24.214	23.217	76.527	1.00 49.49	6
ATOM	1927	CG	LYS A 243	25.061	23.023	77.795	1.00 54.90	6
ATOM	1928	CD	LYS A 243	24.652	23.934	78.939	1.00 58.95	6
ATOM	1929	CE	_	24.782	25.399	78.577	1.00 64.13	
	1930	NZ						6
ATOM				24.274	26.283	79.676	1.00 66.93	7
ATOM	1931	C	LYS A 243	23.965	20.767	76.135	1.00 47.06	6
ATOM	1932	0	LYS A 243	24.735	20.113	76.845	1.00 46.39	8
MOTA	1933	N	GLU A 244	22.716	20.380	75.878	1.00 47.51	7
MOTA	1934	CA	GLU A 244	22.172	19.136	76.429	1.00 51.33	· 6
ATOM	1935	CB	GLU A 244	20.650	19.061	76.259	1.00 54.49	6
ATOM	1936	CG		19.843	20.199	76.842	1.00 62.61	6
ATOM	1937	CD		18.360				
	1938				20.089		1.00 65.15	6
MOTA			1 GLU A 244	17.572	20.980	76.888		8
ATOM	1939	OE:		17.986	19.108	75.807	1.00 64.82	8
ATOM	1940	С	GLU A 244	22.745	17.936	75.698	1.00 50.17	6
ATOM	1941	0	GLU A 244	22.866	16.846	76.259	1.00 51.54	8
ATOM	1942	N	VAL A 245	23.104	18.148	74.441	1.00 47.70	7
ATOM	1943	CA	VAL A 245	23.587	17.063	73.611	1.00 45.43	6
ATOM	1944	CB	VAL A 245	22.704	16.980	72.336	1.00 48.47	6
ATOM	1945		1 VAL A 245	23.082	15.765	71.499	1.00 51.87	6
ATOM	1946		2 VAL A 245					
				21.226	16.934	72.731	1.00 45.65	6
ATOM	1947	C	VAL A 245	25.056	17.070	73.185	1.00 43.01	.6
ATOM	1948	0	VAL A 245	25.620	16.005	72.946	1.00 39.28	8
ATOM	. 1949	N	PHE A 246	25.682	18.245	73.109	1.00 40.53	7
MOTA	1950	CA	PHE A 246	27.063	18.321	72.633	1.00 38.56	6
ATOM	1951	CB	PHE A 246	27.023	18.700	71.154	1.00 36.85	6
ATOM	1952	CG	PHE A 246	28.315	18.487	70.415	1.00 36.46	6
ATOM	1953	CD1	PHE A 246	28.749	17.201	70.098	1.00 32.95	6
ATOM	1954	CD2		29.064	19.582	69.967	1.00 35.51	6
	1955	CE1						
ATOM				29.903	17.004	69.337	1.00 33.80	6
MOTA	1956	CE2		30.222	19.397	69.206	1.00 34.46	6
ATOM	1957	CZ	PHE A 246	30.640	18.103	68.889	1.00 35.54	6
ATOM	1958	С	PHE A 246	27.970	19.311	73.3 7 1	1.00 40.83	6
ATOM	1959	0	PHE A 246	27.613	20.478	73.549	1.00 40.32	8
MOTA	1960	N	GLU A 247	29.141	18.839	73.802	1.00 42.54	7
MOTA	1961	CA	GLU A 247	30.128	19.695	74.467	1.00 43.93	6
ATOM	1962	CB	GLU A 247	30.655	19.075	75.770	1.00 45.67	6
ATOM	1963	CG	GLU A 247	29.763	19.243	77.005	1.00 51.63	6
ATOM	1964	CD	GLU A 247	28.478	18.424	76.962	1.00 57.42	6
			GLU A 247			76.952		
ATOM	1965						1.00 62.12	8
ATOM	1966		GLU A 247	28.296	17.557	77.845	1.00 59.43	8
ATOM	1967	С	GLU A 247	31.268	19.839	73.464	1.00 43.62	6
ATOM ·	1968	0	GLU A 247	32.077	18.931	73.294	1.00 44.25	8
ATOM	1969	N	PRO A 248	31.342	20.988	72.780	1.00 43.65	7
ATOM	1970	CD	PRO A 248	30.439	22.143	72.863	1.00 42.73	6
ATOM	1971	CA	PRO A 248	32.371	21.260	71.779	1.00 43.28	6
ATOM	1972	CB	PRO A 248	31.802	22.480	71.042	1.00 43.23	6
		CG	PRO A 248					
ATOM	1973			30.317	22.474	71.415	1.00 43.02	6
ATOM	1974	С	PRO A 248	33.759	21.552	72.331	1.00 43.37	6
ATOM	1975	0	PRO A 248	33.896	22.286	73.305	1.00 45.44	8
ATOM	1976	N	GLU A 249	34.788	20.982	71.710	1.00 42.38	7
ATOM	1977	CA	GLU A 249	36.151	21.263	72.136	1.00 41.56	6
ATOM	1978	CB	GLU A 249	37.148	20.275	71.528	1.00 42.06	6
ATOM	1979	CG	GLU A 249	36.935	18.816	71.887	1.00 44.28	6
ATOM	1980	CD	GLU A 249	38.015	17.908	71.295	1.00 44.79	6
ATOM	1700	٠.		50.013	11.300	233		•
		•	•					

ATOM	198	1 0	E1 GLU A 2	49	38.20	8 17.938	3 70.054	1.00 42.47	8
ATOM	1982	2 01	E2 GLU A 2	49	38.66				
MOTA	1983		GLU A 2		36.44			1.00 40.99	
ATOM	1984		GLU A 2		37.15	0 23.450			
ATOM	1985		VAL A 2		35.87				
ATOM	1986 1987	_	· ·		36.05				-
MOTA MOTA	1988		SI VAL A 2		37.29 37.12				
ATOM	1989		2 VAL A 2		37.48				
ATOM	1990		VAL A 2		34.830				
ATOM	1991	0	VAL A 2	50	34.162				
MOTA	1992	S N	TYR A 2		34.539				
ATOM	1993				33.368	26.183	67.916	1.00 27.07	6
ATOM	1994				32.185				6
MOTA	1995				32.080				6
MOTA MOTA	1996 1997				31.553 31.439				6
ATOM	1998				32.494		69.106 70.696		6
ATOM	1999				32.384		71.193		6 6
ATOM	2000				31.854		70.391	1.00 34.75	6
ATOM	2001	OH	TYR A 2	51	31.743		70.867	1.00 33.52	8
ATOM	2002		TYR A 2		33.570	27.384	66.992	1.00 27.48	6
ATOM	2003	0	TYR A 2		34.167	•	67.366	1.00 24.78	8
MOTA	2004	N	LEU A 25		33.063		65.773	1.00 24.80	7
ATOM.	2005 2006	CA CB	LEU A 25		33.150 33.631		64.815	1.00 23.40	6
MOTA	2007	CG	LEU A 25		35.126	27.810 27.456	63.451 63.385	1.00 21.32	6 6
ATOM	2008		LEU A 25		35.457		64.395	1.00 22.51	6
MOTA	2009	CD	2 LEU A 25	2	35.499	26.999	61.986	1.00 22.07	6
ATOM	2010	С	LEU A 25		31.762	28.959	64.729	1.00 22.56	6
ATOM	2011	0	LEU A 25		30.750	28.266	64.856	1.00 21.99	8
ATOM	2012	N	LEU A 25		31.734	30.277	64.554	1.00 21.01	7
ATOM ATOM	2013 2014	CA CB	LEU A 25 LEU A 25		30.498	31.047	64.461	1.00 18.89	6
ATOM	2015	CG	LEU A 25		30.352 29.198	31.944 32.942	65.695 65.842	1.00 20.05 1.00 21.61	6
MOTA	2016		LEU A 25		27.849	32.220	65.860	1.00 21.61	. 6 6
ATOM	2017		LEU A 25		29.395	33.716	67.145	1.00 22.90	6
ATOM	2018	С	LEU A 25		30.539	31.901	63.198	1.00 20.05	6
ATOM	2019	0	LEU A 25		31.466	32.691	62.987	1.00 18.17	8
ATOM	2020	N	GLN A 25		29.544	31.720	62.340	1.00 19.40	7
MOTA	2021 2022	CA CB	GLN A 25		29.488	32.490	61.115	1.00 18.17	6
ATOM ATOM	2023	CG	GLN A 25		29.017 27.584	31.592 31.713	59.969 59.601	1.00 9.67 1.00 18.43	6 6
ATOM	2024	CD	GLN A 25		27.368	32.766	58.549	1.00 19.97	6
MOTA	2025	OE1	GLN A 25		27.917	32.677	57.450	1.00 22.54	8
MOTA	2026	NE2			26.564	33.769	58.869	1.00 22.89	7
ATOM	2027	Ç	GLN A 254		28.520	33.634	61.444	1.00 19.75	6
ATOM	2028	0	GLN A 254		27.470	33.415	62.060	1.00 18.77	8
MOTA	2029 2030	N CA	LEU A 255		28.905 28.132	34.854	61.067	1.00 23.02	7
ATOM ATOM	2031	CB	LEU A 255		28.963	36.052 36.993	61.369 62.242	1.00 23.77 1.00 26.84	6 6
ATOM	2032	CG	LEU A 255	•	29.226	36.556	63.684	1.00 29.34	6
ATOM	2033		LEU A 255		30.196	37.520	64.331	1.00 30.65	6
ATOM	2034	CD2	LEU A 255	i	27.902	36.506	64.456	1.00 28.42	6
ATOM	2035	С	LEU A 255		2 7.605	36.842	60.197	1.00 24.84	6
ATOM	2036	0	LEU A 255		27.774	38.066	60.149	1.00 24.94	8
ATOM	2037	N	GLY A 256		26.969	36.158	59.254	1.00 25.07	7
ATOM	2038	CA	GLY A 256		26.408	36.858	58.117	1.00 26.11	6
ATOM ATOM	2039 2040	С 0	GLY A 256 GLY A 256		25.506 24.742	37.956	58.644	1.00 27.11 1.00 25.67	6 8
ATOM	2041	N	THR A 257		25.599	37.734 39.150	59.584 58.072	1.00 27.85	7
ATOM	2042	CA	THR A 257		24.757	40.244	58.536	1.00 29.28	6
MOTA	2043		THR A 257		25.517	41.597	58.545	1.00 27.98	6
MOTA			THR A 257		26.002	41.895	57.232	1.00 31.91	8
MOTA			THR A 257		26.686	41.541	59.510	1.00 26.45	6
MOTA	2046	C	THR A 257		23.477	40.392	57.722	1.00 28.39	6

ATO:	M 2047 O THR A 257	22.747 41.370 57 879 1 00 20 40
ATO!		37.075 1.00 29.49
ATO		23.192 39.414 56.867 1.00 29 13
ATO	1.51 1. 230	21.977 39.471 56.065 1.00 30 49
		22.004 38.432 54.933 1.00 28.22
ATO	11 250	2,00 20,22
ATOI	NOI N 200	33.110 1.00 29.39
ATON	4 2053 OD2 ASP A 258	30.520 1.00 30.16
ATON		23.019 36.292 54.667 1.00 29.97
ATOM		20.046 39.355 56.826 1.00 30.50
		19.601 39.622 56.249 1 00 20 ==
ATOM	1.0 1. 233	20 650 39 012 50 101
ATOM		21 711 38 338 50 053 1 00 22 -
ATOM	1 2058 CA PRO A 259	
ATOM		1.00 31./3
ATOM		19.705 37.912 59.987 1.00 31 87
ATOM		21.067 38.373 60.333 1.00 31.73
		18.81/ 40 18/ 50 260 1 00 20
ATOM		17.736 40 270 50 845 1 20 00
ATOM	2063 N LEU A 260	33.043 1.00 23.78 N
ATOM	2064 CA LEU A 260	1.00 29.38
ATOM	2065 CB LEU A 260	
ATOM		20.36/ 43.542 59.275 1.00 27.74 6
		/ 1.744 //3 796 60 224 4 00 cm
ATOM		22.794 43 919 50 504 1 00 12.17
ATOM	2068 CD2 LEU A 260	21 211 42 702 61 622 4 62 75 6
ATOM	2069 C LEU A 260	02.055 1.00 10.25 6
ATOM	2070 O LEU A 260	12 20 20 20 10 10 20 09 6
ATOM	2071 N LEU A 261	17.787 42.834 57.410 1.00 29.61 R
ATOM	2072 CA LEU A 261	17.237 44.044 59.223 1.00 29.29 7
		10.097 44.693 58 596 1 00 20 71
ATOM	2073 CB LEU A 261	17 740 45 700 EN EST 4 65 55 5
ATOM	2074 CG LEU A 261	14 406 46 664 10 010
ATOM	2075 CD1 LEU A 261	13 144 45 910 50 003 4 00 20.14 6
ATOM	2076 CD2 LEU A 261	30.003 1.00 24.82 6
ATOM	2077 C LEU A 261	
ATOM	2078 O LEU A 261	16.461 45.322 57.259 1.00 29 50
		15./17 45.198 56.295 1 00 31 67 6
ATOM	2079 N GLU A 262	17 603 AE 000 E7 201
MOTA	2080 CA GLU A 262 .	10 015 46 664 57
ATOM	2081 CB GLU A 252	19 049 47 750 56 979 1 00 31.93 6
MOTA	2082 CG GLU A 262	19 406 40 001
ATOM	2083 CD GLU A 262	1.00 20.32
ATOM	2084 OE1 GLU A 262	10.449 48.687 58.589 1.00 29.76 6
ATOM	2085 OE2 GLU A 262	18.175 47.548 59.029 1.00 30.12 8
		18.661 49.661 59.338 1.00 27.69 8
ATOM	2086 C GLU A 262	
ATOM	2087 O GLU A 262	10 600 46 100 70 70
ATOM	2088 N ASP A 263	10 770 44 405
ATOM	2089 CA ASP A 263	7 20 20 20 20 20 20 20 20 20 20 20 20 20
MOTA	2090 CB ASP A 263	21.227 2.00 39.30 6
ATOM	2091 CG ASP A 263	20.354 42.672 54.641 1.00 38.42 6
ATOM		20.982 41.847 53.538 1.00 39.33 6
		22.064 11.263 53.762 1.00 38.86 8
ATOM	2093 OD2 ASP A 263	20 384 11 770 50 447
MOTA	2094 C ASP A 263	
MOTA	2095 O ASP A 263	17 474 41 066 54 301 1.00 42.83
ATOM	2096 N TYR A 264	1.00 44.22
ATOM		
		16.508 42.357 51.796 1.00 46.67 6
ATOM	2098 CB TYR A 264	16.031 43.149 50.568 1.00 54.61 6
MOTA	2099 CG TYR A 264	16 824 42 020 40 004
MOTA	2100 CD1 TYR A 264	2.00 03.02
ATOM	2101 CE1 TYR A 264	2.00 03.73
ATOM	2102 CD2 TYR A 264	17.230 41.709 47.226 1.00 68.35 6
ATOM		17.882 43.786 48.962 1.00 66.23 6
		18.611 43.606 47.780 1.00 68.63 6
ATOM	2104 CZ TYR A 264	18.279 42.570 46.918 1.00 69.08 6
ATOM	2105 OH TYR A 264	10 000
ATOM	2106 C TYR A 264	2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2
MOTA	2107 O TYR A 264	1, 2, 10, 000 1, 100 43.03
	2108 N LEU A 265	17 000
		17.897 40.400 51.333 1.00 40.37 7
		^{18.051} 38.984 51.016 1.00 38.06 6
	2110 CB LEU A 265	19.474 38.646 50.538 1.00 33.48 6
	2111 CG LEU A 265	19.905 39.211 49.172 1.00 32 43 6
ATOM	2112 CD1 LEU A 265	33.222 -2.272 2.00 32.43
		21.176 38.535 48.703 1.00 26.97 6

3 0001	211								
MOTA	211:		02 LEU A 265		18.828			1.00 34.41	6
ATOM	2114		LEU A 265		17.665	38.131	52.219	1.00 37.41	6
MOTA	211	5 0	LEU A 265		18.125	37.000	52.370	1.00 37.96	8
ATOM	2116	5 N	SER A 266		16.804	38.694	53.066		7
ATOM	2117				16.294			1.00 36.46	
ATOM.									6
					17.263			1.00 37.22	6
ATOM	2119				17.190			1.00 37.41	8
MOTA	2120) C	SER A 266		14.997	38.705	5 4 .6 5 3	1.00 35.55	6
ATOM	2121	. 0	SER A 266		14.889	39.927	54.568	1.00 37.09	8
MOTA	2122	N	LYS A 267		14.018			1.00 34.33	7
ATOM	2123				12.750		55.532	1.00 36.13	
ATOM	2124				11.596				6
•								1.00 36.11	6
MOTA	2125				11.503		53.705		6
ATOM	2126	CD			11.453	38.487	52.869	1.00 38.55	6
ATOM_	2127	CE	LYS A 267		11.369	38.170	51.389	1.00 41.60	6
ATOM	2128	NZ	LYS A 267		11.503	39.413	50.569	1.00 42.07	7.
MOTA	2129		LYS A 267		12.791		57.043	1.00 34.90	6
ATOM	2130		LYS A 267		11.758				
							57.694	1.00 37.18	8
ATOM	2131		PHE A 268		13.998		57.595	1.00 32.82	7
ATOM	2132	CA	PHE A 268	•	14.192		59.016	1.00 31.50	6
ATOM	2133	CB	PHE A 268		15.477	38.337	59.495	1.00 34.05	6
ATOM	2134	CG	PHE A 268		15.379	36.839	59.604	1.00 34.54	6.
MOTA	2135	CD:	1 PHE A 268		16.506		59.940	1.00 35.04	6
ATOM	2136		2 PHE A 268		14.161		59.429	1.00 34.57	6
ATOM	2137		PHE A 268						
					16.423	34.691	60.108	1.00 35.44	6
ATOM	2138	CE	_		14.066	34.784	59.594	1.00 36.00	6
ATOM	2139	CZ	PHE A 268		15.201	34.040	59.936	1.00 34.68	6
MOTA	2140	С	PHE A 268		14.319	40.530	59.190	1.00 30.94	6
ATOM	2141	0	PHE A 268		14.983	41.192	58.394	1.00 30.27	8
ATOM	2142	N	ASN A 269		13.693	41.081	60.222	1.00 32.53	7
ATOM	2143	CA	ASN A 269		13.760	42.527	60.448	1.00 35.83	6
ATOM	2144	CB	ASN A 269		12.344	43.115	60.570		
								1.00 37.23	6
ATOM	2145	CG	ASN A 269		11.478	42.809	-	1.00 40.75	6
MOTA	2146		ASN A 269		11.830	43.148	58.227	1.00 43.88	8
MOTA	2147	ND2			10.335	42.165	59.594	1.00 39.61	7
ATOM	2148	С	ASN A 269		14.553	42.854	61.710	1.00 35.45	6
ATOM	2149	0	ASN A 269		14.095	43.621	62.560	1.00 41.47	8
ATOM	2150	N	LEU A 270		15.747	42.285	61.827	1.00 33.27	7
MOTA	2151	CA	LEU A 270		16.571	42.510	63.004	1.00 30.68	6
ATOM	2152	CB	LEU A 270		17.638	41.431	63.114		
	2153	CG	LEU A 270					1.00 27.41	6
ATOM					17.140	40.002	62.988	1.00 23.76	6
MOTA	2154		LEU A 270		18.222	39.106	63.543	1.00 27.14	6
ATOM	2155		LEU A 270		15.855	39.801	63.772	1.00 28.00	6
ATOM	2156	С	LEU A 270		17.258	43.856	63.033	1.00 32.30	6
MOTA	2157	0	LEU A 270		17.347	44.554	62.017	1.00 36.27	8
ATOM	2158	N	SER A 271		17.749	44.207	64.216	1.00 30.33	7
ATOM	2159	CA	SER A 271		18.465	45.457	64.424	1.00 30.79	6
ATOM	2160	CB	SER A 271		17.816	46.249			
	2161		SER A 271				65.562	1.00 29.53	6
MOTA		OG			17.712	45.471	66.739	1.00 30.43	8
MOTA	2162	C	SER A 271		19.911	45.109	64.768	1.00 33.09	6
MOTA	2163	0	SER A 271		20.194	43.972	65.172	1.00 29.64	8
MOTA	2164	N	ASN A 272		20.821		64.586	1.00 32.36	· 7
MOTA	2165	CA	ASN A 272		22.234	45.846	64.896	1.00 31.65	6
ATOM	2166	CB	ASN A 272		23.036	47.141	64.771	1.00 33.76	6
ATOM	2167	CG	ASN A 272		23.101		63.361		6
						47.658		1.00 37.76	
ATOM	2168		ASN A 272		23.719	48.686	63.100	1.00 36.12	8
ATOM	2169		ASN A 272		22.460	46.952	62.437	1.00 44.79	7
ATOM	2170	С	ASN A 272		22.369	45.333	66.321	1.00 32.61	6
ATOM	2171	0	ASN A 272		22.970	44.283	66.565	1.00 27.95	8
ATOM	2172	N	VAL A 273		21.803	46.091	67.257	1.00 33.22	7
ATOM	2173		VAL A 273		21.839	45.741	68.668	1.00 35.52	6
			VAL A 273		20.928			1.00 37.80	6
ATOM	2174					46.660	69.481		
ATCM	2175		VAL A 273		20.987	46.276	70.964	1.00 39.00	6
	2176		VAL A 273		21.356	48.112	69.275	1.00 38.88	6
MOTA	2177		VAL A 273		21.416	44.300	68.908	1.00 34.26	6
ATOM	2178	0	VAL A 273		22.060	43.580	69.679	1.00 35.96	8
							_		

	ATCM	217	9 N	AL	A A 274		20.32	8 43.88	0 (0)(_
	ATCM	218	0 C		A A 274		19.83				
	ATCM			-	A A 274						
	ATCM						18.57			3 1.00 28.88	1 6
			_		A A 274		20.92			1.00 31.74	ϵ
	ATCM				A A 274		21.32	3 40.63	68.533		
	ATCM				E A 275		21.40				
•	ATOM	218	5 C	A PHI	E A 275		22.46				
	MOTA	218	6 C		E A 275		22.93				
	ATOM	218		G PHE	E A 275		23.93				
	ATCM			חום וח	E A 275						
	ATOM			DT LIM	E A 275		23.59				6
			ט פו	72 PNE	A 2/5		25.21				6
	ATOM		0 61	ST PHE	A 275		24.51	3 39.100	62.586	1.00 32.27	
	MOTA				E A 275		26.149	9 40.828	62.950		
	ATCM				A 275		25.793	39.613			-
	MOTA	2193	3 C	PHE	A 275		23.632	40.999	67.040		-
	ATOM	2194	1 0	PHE	A 275		24.252				6
	ATOM	2195	5 N		A 276		23.908				8
	ATCM	2196		LEU	A 276		24.988				7
	ATOM	2197			A 276	٠					6
	ATOM	2198					25.221				6
				, PEO	A 276		26.415			1.00 36.01	6
	ATCM	2199	CL	T LEU	A 276		26.683	45.391	70.025		6
	ATOM	2200			A 276		26.147				6
	ATCM	2201	. С		A 276		24.682				6
	ATOM	2202	0	LEU	A 276		25.560		70.371	1.00 32.79	
	ATOM	2203	N		A 277			41.273	70.384		8
	ATOM	2204	CA		A 277		23.086			1.00 33.95	7
	ATOM	2205			A 277		21.623		71.505	1.00 36.56	6
	ATOM	2206			A 277		21.343		71.902	1.00 35.76	6
	ATOM	2207			A 277				72.687	1.00 42.31	6
	MOTA	2208		LVC	2 2 7 7		20.743		74.049	1.00 45.72	6
			_		A 277		21.665		74.865	1.00 47.87	6
	ATOM	2209			A 277		21.140	40.378	76.244	1.00 44.36	7
	MOTA	2210			A 277		23.302	38.974	71.092	1.00 37.98	6
	MOTA	2211	0		A 277		23.875	38.179	71.845	1.00 37.54	8
	ATOM	2212	N	ALA	A 278		22.832	38.654	69.886	1.00 37.17	7
	ATOM	2213	CA		A 278		22.952	37.311	69.323	1.00 34.51	
	ATOM	2214	CB	ALA	A 278		22.638	37.341	67.820		6
	ATOM	2215	С		A 278		24.368	36.831		1.00 35.38	6
	ATOM	2216	o		A 278		24.605		69.550	1.00 30.63	6
	ATOM	2217	N		A 279			35.790	70.167	1.00 27.62	8
	ATCM	2218	CA		A 279		25.303		69.049	1.00 29.24	7
	ATOM	2219	CB				26.722	37.347	69.167	1.00 31.48	6
					A 279		27.490	38.558	68.645	1.00 33.25	6
	ATOM	2220	CG		A 279		28.974	38.396	68.663	1.00 39.28	6
	ATOM	2221			A 279		29.578	37.337	68.000	1.00 41.15	6
	MCTA	2222		PHE			29.776	39.328	69.315	1.00 40.66	6
	ATOM	2223		PHE			30.960	37.209	67.987	1.00 44.22	6
	ATOM	2224	CE2	PHE .	A 279		31.153	39.213	69.378	1.00 41.38	6
	ATOM	2225	CZ	PHE .	A 279		31.750	38.152	68.614	1.00 44.52	
	ATOM	2226	С		A 279		27.116	37.043	70.6:1		6
	ATOM	2227	0		A 279		27.627			1.00 31.81	6
	ATCM	2228	N		A 280			35.953	70.935	1.00 27.51	8
	ATOM	2229	CA		A 280		26.860	38.005	71.503	1.00 29.32	7
							27.192	37.851	72.907	1.00 29.26	6
	ATOM	2230	CB		A 280		26.927	39.153	73.660	1.00 30.39	6
	ATOM	2231	CG	ASN A	A 280		27.907	40.245	73.278	1.00 30.68	6
	atom	2232	OD1	ASN A	A 280		29.117	40.030	73.303	1.00 33.34	8
	MOTA	2233	ND2	ASN A	A 280		27.395	41.419	72.931	1.00 27.00	7
	ATOM	2234	С	ASN A	A 280		26.524	36.680	73.616	1.00 30.01	6
	ATOM	2235	0	ASN A			27.167	36.004	74.419		
	ATCM	2236	N	ILE 2			25.252		/3.413 73 33-	1.00 29.58	8
	ATOM	2237	CA	ILE A				36.423	73.335	1.00 30.46	7
	ATOM	2238	CB				24.594	35.291	73.983	1.00 33.71	6
				ILE A			23.107	35.161	73.569	1.00 36.14	6
	ATOM	2239	262	ILE A	781		22.541	33.820	74.032	1.00 36.18	6
	ATOM	2240	CG1	ILE A	281		22.298	36.307	74.177	1.00 33.52	6
	ATCM	2241		ILE A			20.835	36.243	73.834	1.00 37.16	6
	ATOM	2242	С	ILE A			25.330	34.006	73.631	1.00 34.06	6
	ATOM	2243	0	ILE A	281		25.385	33.071	74.437	1.00 31.94	8
	ATOM	2244	N	VAL A			25.896	33.960	72.427	1.00 35.31	7
	•				• •			55.500		7.00 22.37	,

		•				
ATOM	1 2245 CA .VAL A 282	26.65	32.785	72.00	5 1 00 36 45	_
ATOM		27.084				6
ATOM		27.829				6
'ATOM						
ATOM		25.880				
ATOM		27.919				6
		28.182		73.532	2 1.00 36.12	8
MOTA		28.693		72.821	1.00 38.45	
ATOM		29.929	33.884			6
MOTA		30.551	35.272			
MOTA	2254 CG ARG A 283	30.974	35.625	72.027		6
MOTA	2255 CD ARG A 283	31.492		71.968		6
ATOM		32.647				6
ATOM		33.162		72.840		7
ATOM	2258 NH1 ARG A 283	32.628		73.215		6
ATOM	2259 NH2 ARG A 283			72.797		7
MOTA	2260 C ARG A 283	34.220		74.014		7
ATOM		29.614		75.044		6
ATOM		30.350		75.716	1.00 39.01	8
		28.506	34.141	75.520	1.00 40.30	7
MOTA	2263 CA GLU A 284	28.084	33.923	76.894	1.00 43.19	6
ATOM	2264 CB GLU A 284	26.753	34.647	77.165		6
MOTA	2265 CG GLU A 284	26.875	36.176	77.090	1.00 56.10	6
MOTA	2266 CD GLU A 284	25.542	36.923	77.179	1.00 60.77	6
ATOM	2267 OE1 GLU A 284	24.659	36.682	76.329	1.00 61.41	
ATOM	2268 OE2 GLU A 284	25.383	37.763	78.096	1.00 62.21	8
ATOM	2269 C GLU A 284	27.953	32.429	77.179	1.00 40.72	8
ATOM	2270 O GLU A 284	28.565	31.922	78.120		6
ATOM	2271 N VAL A 285	27.186	31.721	76.354	1.00 45.29	8
ATOM	2272 CA VAL A 285	26.975	30.288	76.551	1.00 34.82	7
ATOM	2273 CB VAL A 285	25.842			1.00 30.84	6
ATOM	2274 CG1 VAL A 285	25.698	29.752 28.253	75.647	1.00 27.74	6
ATOM	2275 CG2 VAL A 285	24.545		75.831	1.00 22.95	6
ATOM	2276 C VAL A 285		30.433	75.982	1.00 26.26	6
ATOM	2277 O VAL A 285	28.181	29.366	76.341	1.00 31.93	6
ATOM		28.492	28.556	77.214	1.00 33.46	8
ATOM		28.845	29.466	75.191	1.00 29.43	7
		29.973	28.586	74.907	1.00 24.26	6
ATOM	2280 CB PHE A 286	29.830	27.957	73.519	1.00 22.57	6
ATOM	2281 CG PHE A 286	28.607	27.095	73.345	1.00 23.46	6
MOTA	2282 CD1 PHE A 286	27.409	27.639	72.885	1.00 23.90	6
ATOM	2283 CD2 PHE A 286	28.664	25.718	73.608	1.00 21.95	6
ATOM	. 2284 CE1 PHE A 286	26.281	26.814	72.681	1.00 24.90	6
ATOM	2285 CE2 PHE A 286	27.547	24.892	73.411	1.00 18.06	6
MOTA	2286 CZ PHE A 286	26.357	25.437	72.945	1.00 20.23	6
ATOM	2287 C PHE A 286	31.368	29.200	74.991	1.00 25.14	6
ATOM	2288 O PHE A 286	32.338	28.566	74.560	1.00 23.14	8
MOTA	2289 N GLY A 287	31.480	30.416	75.525	1.00 25.51	
ATOM	229· CA GLY A 287	32.783	31.065	75.614		7
ATOM	2291 C GLY A 287	33.353		74.270	1.00 26.86	6
ATOM	2292 O GLY A 287	32.644	31.549		1.00 26.28	6
MOTA	2293 N. GLU A 288	34.637		73.271	1.00 26.29	8
ATOM	2294 CA GLU A 288		31.849	74.238	1.00 27.17	7
MOTA	2295 CB GLU A 288	35.274		72.996	1.00 33.20	6
ATOM	2296 CG GLU A 288	36.680		73.269	1.00 35.09	6
		36.726	34.104		1.00 41.67	6
MOTA		35.970	35.231	73.421	1.00 43.13	6
MOTA	2298 OE1 GLU A 288	36.221	35.493	72.228	1.00 45.39	8
MOTA	2299 OE2 GLU A 288	35.130	35.858	74.097	1.00 47.64	8
MOTA	2300 C GLU A 288	35.386	31.204	71.930	1.00 32.87	6
MOTA	2301 O GLU A 288	35.596		72.247	1.00 31.59	8
MOTA	2302 N GLY A 289			70.668	1.00 31.93	7
MOTA	2303 CA GLY A 289			69.545	1.00 29.58	6
MOT	2304 C GLY A 289		_	68.307	1.00 27.42	5
MOTA	2305 O GLY A 289			68.398	1.00 27.42	
TOM	2306 N VAL A 290					8
TOM	2307 CA VAL A 290			67.143	1.00 27.64	7
TOM	2308 CB VAL A 290				1.00 27.17	6
TOM	2309 CG1 VAL A 290			55.037	1.00 25.25	6
					1.00 22.41	6
TOPI	2310 CG2 VAL A 290		29.632 [55.852	1.00 17.33	6
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ATOM	2311	С	VAL A	290	35.137	31.975	65.105	1.00 25.97	6
ATOM	2312	0	VAL A	290	34.218	31.279	64.672	1.00 22.32	8
		N	TYR A			33.293	64.914		
MOTA	2313				35.217			1.00 27.33	7
MOTA	2314	CA		-	34.188	34.052	64.203	1.00 26.69	6
ATOM	2315	CB	TYR A	291	33.925	35.356	64.939	1.00 25.51	6
ATOM	2316	CG	TYR A	291	33.935	35.178	66.435	1.00 28.73	6
ATOM	2317		1 TYR A		35.025	35.596	67.191	1.00 29.72	6
	2318		1 TYR A			35.414	68.563		
MOTA		-			35.059			1.00 29.53	6
MOTA	2319	CD	2 TYR A		32.874	34.565	67:094	1.00 27.39	6
MOTA	2320	CE.	2 TYR A 2	291	32.898	34.377	68.466	1.00 31.03	6
MOTA	2321	CZ	TYR A 2	291 -	33.997	34.808	69.194	1.00 31.85	6
MOTA	2322	ОН	TYR A	291	34.030	34.647	70.562	1.00 38.03	8
	2323	C	TYR A		34.527	34.345	62.745	1.00 27.99	6
MOTA									
ATOM	2324	.0	TYR A 2		35.608	34.843	62.415	1.00 24.83	8
ATOM	2325	N	LEU A 2		33.567	34.042	61.880	1.00 30.17	7
ATOM	2326	CA	LEU A 2		33.726	34.220	60.441	1.00 28.26	6
ATOM	2327	CB	LEU A 2	292	. 33,561	32.861	59.741	1.00 27.70	6
ATOM	2328	CG	LEU A 2	292	34.191	31.643	60.435	1.00 24.64	6
ATOM	2329	CD	L LEU A 2	292	33.867	30.380	59.661	1.00 24.66	6
ATOM	2330		LEU A 2		35.686	31.825	60.553	1.00 23.19	
	2331	c c	LEU A 2		32.649	35.175	59.944	1.00 25.59	6
ATOM			LEU A 2						
ATOM	2332	0			31.640	35.394	60.611	1.00 18.11	8
ATOM	2333	N	GLY A 2		32.869	35.749	58.770	1.00 28.55	7
ATOM	2334	CA	GLY A 2	.93	31.878	36.653	58.223	1.00 31.26	6
ATOM'	2335	С	GLY A 2	93	30.722	35.815	57.714	1.00 34.84	6
ATOM	2336	0	GLY A 2	93	30.463	34.724	58.234	1.00 34.11	8
ATOM	2337	N	GLY A 2	94	30.036	36.312	56.689	1.00 35.34	7
ATOM	2338	CA	GLY A 2		28.918	35.581	56.124	1.00 34.84	6
	2339	c c	GLY A 2		28.142	36.445	55.155	1.00 34.79	6
MOTA		ō	GLY A 2				54.699	1.00 37.05	8
MOTA	2340			-	28.644	37.473			
MOTA	2341	N	GLY A 2		26.917	36.035	54.842	1.00 31.87	7
MOTA	2342	CA	GLY A 2		26.102	36.806	53.925	1.00 27.78	6
ATOM	2343	С	GLY A 2		25:969	38.245	54.378	1.00 27.09	6
ATOM	2344	, О	GLY A 2	95	26.192	38.558	55.546	1.00 27.03	8
ATOM	2345	N	GLY A 2	96	25.596	39.119	53.450	1.00 24.67	7
ATOM	2346	CA	GLY A 2	96	25.440	40.527	53.757	1.00 25.28	6
ATOM	2347	C	GLY A 2		25.562	41.262	52.446	1.00 27.64	6
	2348	ŏ	GLY A 2		26.591	41.163	51.771	1.00 26.65	8
ATOM	2349		TYR A 2		24.526		52.078	1.00 30.21	7
ATOM		Ŋ				42.009			
ATOM	2350	CA	TYR A 2		24.543	42.704	50.801	1.00 30.62	6
ATOM	2351	CB	TYR A 2		23.560	42.011	49.859	1.00 29.50	6
ATOM	2352	CG	TYR A 2		23.717	40.516	49.953	1.00 30.33	6
ATOM	2353	CD1	TYR A 2	97	23.174	39.810	51.031	1.00 30.86	6
MOTA	2354	CE1	TYR A 2	97	23.450	38.449	51.226	1.00 30.74	6
ATOM	2355	CD2	TYR A 2	97	24.538	39.824	49.062	1.00 31.20	6
MOTA	2356	CE2	TYR A 2	97	24.821	38.460	49.247	1.00 32 08	6
	2357	cz	TYR A 2		24.275	37.781	50.332	1.00 30.92	6
MOTA		ОН	TYR A 2		24.539		50.509	1.00 29 60	8
ATOM	2358					36.440			6
MOTA	2359	C	TYR A 2		24.267	44.195	50.875	1.00 32.07	
ATOM	2360	0	TYR A 2		24.134	44.849	49.840	1.00 33.83	8
ATOM	2361	N	HIS A 2		24.180	44.725	52.094		7
ATOM	2362	CA	HIS A 2	98	23.961	46.153	52.289	1.00 33.94	6
ATOM	2363	CB	HIS A 2	98	22.761	46.430	53.194	1.00 34.75	6
MOTA	2364	CG	HIS A 2	98	22.379	47.880	53.256	1.00 35.16	6
	2365		HIS A 2		22.558	48.809	54.224	1.00 35.72	6
MOTA			HIS A 2		21.779	48.538	52.205	1.00 34.10	7
MOTA	2366						52.522		6
ATOM	2367		HIS A 2		21.605	49.809		1.00 31.84	
ATOM	2368		HIS A 2		22.069	50.000	53.742	1.00 35.46	7
ATOM	2369	С	HIS A 2		25.213	46.697	52.962	1.00 36.21	6
ATOM	2370	0	HIS A 2	98	25.471	46.405	54.133	1.00 33.83	8
ATOM	2371	N	PRO A 25	99	25.992	47.519	52.234	1.00 36.69	7
ATOM	2372	CD	PRO A 25	99	25.680	47.997	50.881	1.00 35.57	6
	2373	CA	PRO A 25		27.238	48.142	52,689	1.00 36.17	6
ATOM		CB	PRO A 25		27.586	49.073	51.525	1.00 37.75	6
MOTA	2374	CG	PRO A 25		26.216	49.399	50.954	1:00 37.76	6
ATOM	2375		PRO A 25				54.000	1.00 34.47	6
MOTA	2376	С	PRU A 23		27.045	48.886	24.000	4.00 34.4/	•
		•		•					

MOTA	1 2377 O PRO A 29	3	27.78	1 48.67	0 54 06		
MOTA			26.05				
ATOM	-						
ATOM			25.74				' 6
			24.496			9 1.00 35.56	6
ATOM			24.648		4 54.02	B 1.00 35.96	
ATOM			25.370	52.36	7 52.843		
ATOM			25.461	53.40	5 51.90		
ATOM			24.016	53.752			
MOTĂ	2385 CE2 TYR A 300		24.098	54.793			
ATOM	2386 CZ TYR A 300		24.823				
ATOM	2387 OH TYR A 300		24.927			1.00 38.09	
ATOM			25.497				
ATOM		-	26.062				
ATOM			24.661			1.00 30.62	
ATOM							7
ATOM			24.323				6
ATOM			23.216				6
			25.539				6
ATOM			25.848			1.00 31.91	8
ATOM			26.223	46.192	56.557		7
ATOM	2396 CA LEU A 302		27.404	45.383	56.795	1.00 30.55	6
ATOM	2397 CB LEU A 302		28.012	45:002	55.441		6
ATOM	2398 CG LEU A 302	. :	29.315	44.223			6
ATOM	2399 CD1 LEU A 302		29.491	43.781			6
ATOM	2400 CD2 LEU A 302		30.475			1.00 32.23	6
ATOM	2401 C LEU A 302		28.418	46.136			
ATOM	2402 O LEU A 302	. 2	8.796	45.676			6
ATOM	2403 N ALA A 303		8.842	47.299			8
ATOM	2404 CA ALA A 303		9.818	48.119	57.173		7
ATOM	2405 CB ALA A 303		0.026	49.424	57.137		6
ATOM	2406 C ALA A 303		9.397	48.397	59.305		6
ATOM	2407 O ALA A 303		0.088			1.00 25.06	6
ATOM	2408 N ARG A 304		8.258	48.015	60.248	1.00 26.90	8
ATOM	2409 CA ARG A 304		-	49.054	59.472	1.00 24.06	7
ATOM	2410 CB ARG A 304		7.794	49.382	60.810	1.00 24.37	6
MOTA	2411 CG ARG A 304	_	6.420	50.052	60.758	1.00 23.99	6
ATOM	2411 CG ARG A 304 2412 CD ARG A 304		6.328	51.257	59.815	1.00 28.77	6
ATOM	2412 CD ARG A 304 2413 NE ARG A 304		5.106	52.089	60.156	1.00 29.96	6
			3.943	51.233	60.369	1.00 36.43	7
ATOM	_		2.893	5 1.573	61.110	1.00 37.01	6
ATOM	2415 NH1 ARG A 304		2.854	52.757	61.713	1.00 36.37	7
ATOM	2416 NH2 ARG A 304		1.896	50.719	61.269	1.00 34.36	7
MOTA	2417 C ARG A 304		7.727	48.142	61.691	1.00 24.24	6
MOTA	2418 O ARG A 304		8.343	48.099	62.762	1.00 22.34	8
ATOM	2419 N ALA A 305	20	5.994	47.132	61.221	1.00 24.51	7
ATOM	2420 CA ALA A 305	20	5.801	45.883	61.959	1.00 22.70	6
ATOM	2421 CB ALA A 305	25	5.880	44.960	61.175	1.00 18.13	6 .
MOTA	2422 C ALA : 305	28	3.089	45.142	62.351	1.00 23.33	. 6
ATOM	2423 O ALA . 305	28	3.237	44.725	63.506	1.00 21.51	8
MOTA	2424 N TRP 306	29	.016	44.961	61.411	1.00 22.79	7
ATOM	2425 CA TRP A 306		.244	44.270	61.764	1.00 24.33	6
MOTA	2426 CB TRP A 306	31	.029	43.842	60.524	1.00 26.93	6
ATOM	2427 CG TRP A 306					1.00 27.96	6
ATOM	2428 CD2 TRP A 306		.861	42.013	58.629	1.00 26.38	
ATOM	2429 CE2 TRP A 306		.366	40.688	58.570	1.00 24.96	6
ATOM	2430 CE3 TRP A 306	31	.462	42.563			6.
ATOM	2431 CD1 TRP A 306		.983		57.490	1.00 23.00	6
ATOM	2432 NE1 TRP A 306			41.484	60.620	1.00 28.53	6
ATOM	2433 CZ2 TRP A 306			40.392	59.797	1.00 25.62	7
ATOM	2434 CZ3 TRP A 306		.450	39.904	57.414	1.00 24.51	6
		31		41.784	56.343	1.00 25.37	6
MOTA	2435 CH2 TRP A 306			40.465	56.315	1.00 24.20	6
MOTA	2436 C TRP A 306			45.108	62.676	1.00 26.51	6
ATOM	2437 O TRP A 306			44.570	63.464	1.00 25.07	8
MOTA	2438 N THR A 307			46.427	62.575	1.00 28.08	7
MOTA	2439 CA THR A 307		.785	47.323	63.415	1.00 27.91	6 .
	2440 CB THR A 307		484	48.796	63.100	1.00 27.86	6
MOTA	2441 OG1 THR A 307		.994 .	49.119	61.799	1.00 30.17	8
MOTA	2442 CG2 THR A 307	32.		49.704	64.137	1.00 24.72	6
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Figure 17-38

		_						
MOTA .	244	3 C	THR A 307	31.44:	1 47.041	64.863	1.00 29.35	6
MOTA	244	4 0	THR A 307	32.31			. ~ 25.55	
MOTA	244	5 N		30.159				8
MOTA				29.740				
ATOM								6
				28.256				6
ATOM	244			27.338		66.058	1.00 34.79	6
MOTA			D1 LEU A 308	25.903	46.887	66.153		
MOTA	2450) C	D2 LEU A 308	27.569				
ATOM	2453	L C	LEU A 308	30.531				
ATOM	2452		LEU A 308	31.230				6
ATOM	2453		ILE A 309					
ATOM	2454			. 30.423				7
				31.108				6
ATOM	2455			30.939		65.431	1.00 34.95	6
ATOM	2456		32 ILE A 309	31.733		65.799	1.00 31.21	6
ATOM	2457	C	31 ILE A 309	29.445	41.631	65.212		6
MOTA	2458	CI		28.726				6
ATOM	2459	C	ILE A 309	32.589		66.772		
ATOM	2460		ILE A 309	33.183	42.617			6
ATOM	2461		TRP A 310			67.657		8
MOTA	2462			33.197		65.977		. 7
				34.612		66.169	1.00 35.26	6
ATOM	2463			35.150	45.311	65.075	1.00 32.61	6
ATOM	2464			36.619	45.588	65.220	1.00 30.79	6
ATOM	2465	CE		37.679	44.620	65.274	1.00 29.93	6
MOTA	2466	CE	2 TRP A 310	38.882	45.330	65.474	1.00 28.42	6
ATOM	2467	CE	3 TRP A 310	37.731	43.224	65:174	1.00 31.59	
MOTA	2468	CD		37.206	46.804	65.380		6
ATOM	2469			38.565	46.659		1.00 30.62	6
ATOM	2470	CZ		40.126		65.536	1.00 29.37	7
ATOM	2471	CZ			44.691	65.578	1.00 27.91	6
	2472	CH		38.978	42.585	65.279	1.00 28.06	6
ATOM-				40.150	43.322	65.479	1.00 26.50	6
ATOM	2473	C	TRP A 310	34.744	45.040	67.545	1.00 36.00	6
ATOM	2474	0	TRP A 310	35.365	44.476	68.440	1.00 36.24	8
MOTA	2475	N	CYS A 311	34.134	46.213	67.715	1.00 34.57	7
MOTA	2476	ÇA	CYS A 311	34.183	46.937	68.985	1.00 32.82	6
ATOM	2477	CB	CYS A 311	33.169	48.085	68.996	1.00 35.62	6
ATOM	2478	SG	CYS A 311	33.439	49.401	67.796	1.00 32.36	16
ATOM	2479	С	CYS A 311	33.912	46.061	70.206	1.00 32.30	
ATOM	2480	0	CYS A 311	34.452	46.313	71.280		6
ATOM	2481	N	GLU A 312	33.062	45.049		1.00 29.82	8
ATOM	2482	CA	GLU A 312			70.053	1.00 32.57	7
ATOM	2483	CB	GLU A 312	32.731	44.159	71.171	1.00 33.86	6
				31.557	43.252	70.807	1.00 34.19	6
ATCM	2484	CG	GLU A 312	30.442	43.185	71.844	1.00 40.27	6
ATOM	2485	CD	GLU A 312	30.923	42.821	73.239	1.00 43.80	6
ATOM	2486	OE:		31.685	41.831	73.383	1.00 44.81	8
MOTA	2487	OE2		30.516	43.522	74.195	1.00 41.54	8
ATOM	2488	C	GLU A 312	33.953	43.298	71.456	1.00 33.77	6
ATOM	2489	0	GLU A 312	34.253	42.957	72.603	1.00 32.07	8
ATOM	2490	N	LEU A 313	34.647	42.945	70.382	1.00 33.45	7
ATOM	2491	CA	LEU A 313	35.848	42.135	70.473	1.00 33.43	
ATOM	2492	CB	LEU A 313	36.172	41.513			6
ATOM	2493	CG	LEU A 313			69.115	1.00 32.14	6
				35.154	40.493	68.626	1.00 27.73	6
ATCM	2494	CDI	LEU A 313	35.587	39.956	67.269	.1.00 30.39	6
ATCM	2495		LEU A 313	35.053	39.367	69.648	1.00 27.87	6
ATOM	2496	С	LEU A 313	36.976	43.031 -	70.903	1.00 31.64	6
MOTA	2497	0	LEU A 313	37.605	42.793	71.925	1.00 31.74	8
ATCM	2498	N	SER A 314	37.206	44.064	70.099	1.00 33.49	7
ATOM	2499	CA	SER A 314	38.232	45.067	70.328	1.00 35.59	6
ATOM	2500	CB	SER A 314	38.107	46.154	69.256	1.00 36.47	
ATOM	2501	OG	SER A 314					6
				39.141	47.120	69.353	1.00 44.55	8
ATCM	2502	C	SER A 314	38.046	45.661	71.730	1.30 37.82	6
ATOM	2503	0	SER A 314	39.015	46.043	72.395	1.00 37.32	8
ATCM	2504	N	GLY A 315	36.794	45.725	72.175	1.00 38.05	7
ATCM	2505	CA	GLY A 315	36.506	46.243	73.498	1.00 42.42	6
ATOM	2506	C	GLY A 315	36.295	47.744	73.568	1.00 46.80	6
ATOM	2507	0	GLY A 315	35.923	48.276	74.618	1.00 47.85	8
ATCM	2508	N	ARG A 316	36.518	48.438	72.458	1.00 48.90	7.
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ATOM	2509 CA ARG A 316	36.346 49.88	35 72.448 1.00 52.27	_
ATOM	2510 CB ARG A 316	37.144 50.41	70 71 000 1 00 7	6
ATOM		36.730 50.00	07 69.900 1.00 52.11	6
ATOM ATOM	510	37.734 50.51	l4 68.870 1.00 53.76	6
ATOM		39.028 49.85 40.135 50.22	1 60 000	7
MOTA	2515 NH1 ARG A 316	40.135 50.22 40.110 51.25		6
MOTA	2516 NH2 ARG A 316	41.266 49.54		7 7
MOTA		34.882 50.34	2 22 22	, 6
ATOM		34.075 49.78	1 71.652 1.00 54.96	8
MOTA MOTA		34.547 51.36		7
MOTA ·	2521 CB GLU A 317	33.185 51.90 33.111 53.13	0 74 400	6
ATOM	2522 CG GLU A 317	32.549 52.90	1 77 707	5
ATOM	2523 CD GLU A 317	33.353 51.91		
ATOM	2524 OE1 GLU A 317	33.025 51.74	1 77.556 1.00 64.59 8	
ATOM ATOM	2525 OE2 GLU A 317 2526 C GLU A 317	34.305 51.30	2 75.832 1.00 68.64 8	3
ATOM	2527 0 GLU A 317	32.642 52.25 33.270 52.98		
ATOM	2528 N VAL A 318	33.270 52.98 31.457 51.73		
ATOM	2529 CA VAL A 318	30.780 51.96		
MOTA	2530 CB VAL A 318	29.522 51.07		
MOTA	. 2531 CG1 VAL A 318 2532 CG2 VAL A 318	28.875 51.23	7 68.808 1.00 45.53 6	
ATOM ATOM	2533 C VAL A 318	29.895 49.633 30.349 53.413		
ATOM	2534 O VAL A 318	30.349 53.413 29.511 53.867		
ATOM	2535 N PRO A 319	30.925 54.169		
ATOM	2536 CD PRO A 319	31.960 53.836	68.247 1.00 48.87 6	
ATOM ATOM	2537 CA PRO A 319 2538 CB PRO A 319	30.538 55.569	69.093 1.00 52.54 6	
ATOM	2539 CG PRO A 319	31.438 56.051 31.612 54.802	· · · · · · · · · · · · · · · · · · ·	
ATOM	2540 C PRO A 319	31.612 54.802 29.052 55.679		
MOTA	2541 O PRO A 319	28.531 54.913		
ATOM	2542 N GLU A 320	28.369 56.624		
ATOM ATOM	2543 CA GLU A 320 2544 CB GLU A 320	26.942 56.804		
ATOM	2545 CG GLU A 320	26.302 57.588 26.727 59.042		
ATOM	2546 CD GLU A 320	26.007 59.823		
ATOM	2547 OE1 GLU A 320	24.755 59.832		
ATOM ATOM	2548 OE2 GLU A 320 2549 C GLU A 320	26.697 60.431	72.303 1.00 79.46 8	
ATOM	2549 C GLU A 320 2550 O GLU A 320	26.698 57.551 25.663 58.197	67.863 1.00 61.40 6	
ATOM	2551 N LYS A 321	25.663 58.197 27.650 57.463	67.699 1.00 62.33 8 66.939 1.00 59.47 7	
ATOM	2552 CA LYS A 321	27.519 58.150	66.939 1.00 59.47 7 65.662 1.00 59.54 6	
MOTA	2553 CB LYS A 321	27.340 59.648	65.897 1.00 61.36 6	
ATOM ATOM	2554 CG LYS A 321 2555 CD LYS A 321	23.620 60.323	66.366 1.00 65.23 6	
ATOM	2555 CD LYS A 321 2556 CE LYS A 321	13.169 59.691 33.564 60.215	67.643 1.00 66.59 6	
ATOM	2557 NZ LYS A 321	33.564 60.215 30.591 61.699	67.960 1.00 67.34 6 68.100 1.00 68.58 7	
ATOM	2558 C LYS A 321	28.766 57.941	68.100 1.00 68.58 7 64.806 1.00 59.24 6	
ATOM	2559 O LYS A 321	29.845 57.623	65.319 1.00 58.70 8	
ATOM	2560 N LEU A 322	28.608 58.146	63.500 1.00 57.55 7	
ATOM ATOM	2561 CA LEU A 322 2562 CB LEU A 322	29.702 58.002	62.543 1.00 54.72 6	
ATOM	2563 CG LEU A 322	29.171 57.450 28.141 56.316	61.214 1.00 52.96 6 61.295 1.00 52.97 6	
ATOM	2564 CD1 LEU A 322	27.708 55.932	61.295 1.00 52.97 6 59.899 1.00 49.01 6	
MOTA	2565 CD2 LEU A 322	28.716 55.115	62.035 1.00 54.73 6	
MOTA	2566 C LEU A 322	30.250 59.406	62.313 1.00 53.47 .6	
MOTA MOTA	2567 O LEU A 322 2568 N ASN A 323	29.512 60.383	62.464 1.00 53.39 8	
ATOM		31.530 59.521 32.089 60.842	61.965 1.00 51.43 7	
ATOM	2570 CB ASN A 323	33.591 60.905	61.706 1.00 50.32 6 62.035 1.00 52.31 6	
MOTA	2571 CG ASN A 323	34.428 59.964	61.189 1.00 55.06 6	
MOTA	2572 OD1 ASN A 323	34.386 58.744	61.363 1.00 55.77 8	
MOTA	2573 ND2 ASN A 323	35.195 60.530	60.259 1.00 52.71 7	
NTOM	2574 C ASN A 323	31.843 61.199	60.243 1.00 48.63 6	

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ATOM				323	31.13	60.479	59.538	1.00 47.52	8
ATOM	257				32.426	62.304	59.792		
ATOM	257				32.242				
MOTA MOTA	2579 2579				32.758			1.00 50.73	_
ATOM	2586		g asna 3 D1 asna 3		32.025			1.00 53.83	_
ATOM	2583		D2 ASN A 3		30.812 32.755			1.00 56.90	
ATOM	2582		ASN A 3		32.906			1.00 54.93 1.00 49.34	
ATOM	2583		ASN A 3		32.275			1.00 47.22	
ATOM	2584	1 N	LYS A 3		34.182			1.00 48.27	
ATOM	2585	5 C			34.957	60.759	56.676	1.00 46:57	
ATOM	2586				36.314		57.305	1.00 49.94	
MOTA	2587				37.299		56.399	1.00 54.75	
MOTA	2588				38.562		57.173	1.00 58.22	6
ATOM ATOM	2589 2590				39.236		57.844	1.00 58.35	6
ATOM	2591		LYS A 3	-	40.473 34.202		58.566	1.00 59.81	7
ATOM	2592		LYS A 3		34.202		56.410 55.263	1.00 44.85	6
ATOM	2593		ALA A 3		33.712		57.483	1.00 43.59 1.00 42.27	. 8
ATOM	2594				32.964	57.597	57.387	1.00 42.27	7 6
MOTA	2595	CE			32.663	57.067	58.773	1.00 36.86	6
MOTA	2596		ALA A 3	26	31.666	57.803	56.612	1.00 43.30	6
ATOM	2597		ALA A 3		31.342	57.028	55.705	1.00 42.83	8
ATOM	2598		LYS A 3		30.918	58.843	56.977	1.00 45.56	7
ATOM ATOM	2599 2600				29.657	59.146	56.306	1.00 47.23	6
ATOM	2601				29.023 28.547	60.407	56.892	1.00 49.59	6
ATOM	2602	CD			28.024	60.263 61.591	58.329 58.862	1.00 54.63	6
ATOM	2603	CE			27.529	61.483	60.299	1.00 55.89 1.00 58.28	6 6
ATOM	2604	NZ	LYS A 32	27	26.304	60.644	60.426	1.00 59.91	7
ATOM	2605	С	LYS A 32		29.888	59.347	54.816	1.00 46.97	6
MOTA	2606	0	LYS A 32		29.090	58.913	53.990	1.00 48.10	8
ATOM	2607	N	GLU A 32			60.012	54.480	1.00 44.99	7
ATOM ATOM	2608 .2609	CA CB	GLU A 32 GLU A 32		31.325	60.264	53.091	1.00 43.18	6
MOTA	2610	CG	GLU A 32		32.417 [/] 31.993	61.326 62.621	53.027 53.710	1.00 47.93	6
ATOM	2611	CD	GLU A 32		33.112	63.630	53.710	1.00 53.65 1.00 55.79	6 6
ATOM	2612	OE:			33.642	64.060	52.783	1.00 58.73	8
MOTA	2613	OE:	2 GLU A 32		33.459	63.991	54.979	1.00 58.13	8
ATOM	. 2614	С	GLU A 32		31.789	58.971	52.437	1.00 41.56	6
ATOM	2615	0	GLU A 32		31.537	58.743	51.255	1.00 39.41	8
ATOM ATOM	2616 2617	N CA	LEU A 32 LEU A 32		32.465	58.123	53.211	1.00 40.64	7
ATOM	2618	CB	LEU A 32		32.940 33.623	56.844	52.695	1.00 36.45	6
ATOM	2619	CG	LEU A 32		34.100	56.032 54.610	53.801 53.433	1.00 34.70 1.00 35.69	6
ATOM	2620		LEU A 32		35.195	54.642	52.359	1.00 30.11	6 6
ATOM	2621	CD2	LEU A 32	9	34.619	53.926	54.683	1.00 34.63	6
MOTA	2622	С	LEU A 32		31.746	56.064	52.157	1.00 35.77	6
MOTA	2623	0	LEU A 32		31.692	55.746	50.975	1.00 34.94	8
ATOM	2624		LEU A 33		30.784	55.770	53.029	1.00 34.78	7
ATOM	2625 2626		LEU A 33	_	29.599			1.00 34.95	6
ATOM ATOM	2627	CB CG	LEU A 33		28.631 29.164	54.914	53.803	1.00 30.95	6
ATOM	2628		LEU A 33		28.051	54.115 53.904	54.991 56.022	1.00 32.66 1.00 31.74	6
ATOM	2629		LEU A 33		29.674	52.769	54.509	1.00 31.74	6 6
MOTA	2630	С	LEU A 330		28.877	55.631	51.428	1.00 37.28	6
ATOM	2631	0	LEU A 330		28.395	54.901	50.557	1.00 40.56	8
MOTA	2632	N	LYS A 33:		28.806.	56.957	51.383	1.00 38.24	7
ATOM	2633	CA	LYS A 331		28.140	57.661	50.294	1.00 39.59	6
ATOM	2634	CB	LYS A 331		27.994	59.146	50.643	1.00 42.31	C
ATOM	2635	CG	LYS A 331		27.129	59.399	51.873	1.00 45.93	6
MOTA	2636	CD	LYS A 331		27.017	60.879	52.244	1.00 49.72	6
ATOM ATOM	2637 2638	CE NZ	LYS A 331		26.271 26.053	61.698	51.193	1.00 53.66	6
ATOM	2639	C	LYS A 331		28.863	63.114 57.514	51.640 48.958	1.00 54.22 1.00 41.02	7 6
ATOM	2640	ō	LYS A 331		28.220	57.485	47.904	1.00 41.02	8
						J , , 400			-

MOTA	2641	. N	SER A 332	30.192	57.413	49.005	1.00 42.69	7
ATOM	2642	: C		30.998		47.792	1.00 46.02	
								6
MOTA	2643			32.494		48.124	1.00 47.51	6
MOTA	2644	00	SER A 332	32.862	56.066	48.823	1.00 50.97	8
ATOM	2645	C	SER A 332	30.634	56.007	47.040	1.00 48.51	6
ATOM	2646		SER A 332	30.706		45.811		
							1.00 49.45	8
MOTA	2647		ILE A 333	30.241	54.982	47.786	1.00 51.56	7
ATOM	2648	CZ	ILE A 333	29.869	53.713	47.187	1,00 54.86	6
MOTA	2649	CE	ILE A-333	29.657	52.626	48.246	1.00 55.80	6
	2650		2 ILE A 333					
ATOM				29.388		47.559	1.00 52.34	6
MOTA	2651	CG	1 ILE A 333	30.892	52.522	49.140	1.00 56.89	6
MOTA	2652	CI	1 ILE A 333	30.766	51.456	50.204	1.00 60.31	6
ATOM	2653	С	ILE A 333	28.579	53.813	46.396	1.00 57.07	6
	2654		ILE A 333	27.572				
MOTA					54.321	46.897	1.00 55.59	8
MOTA	2655		ASP A 334	28.623	53.320	45.160	1.00 61.14	7
MOTA	2656	CA	ASP A 334	27.456	53.300	44.281	1.00 65.55	6
ATOM	2657	CB	ASP A 334	27.888	53.259	42.811	1.00 69.19	6
	2658			28.784	52.073	42.491		
MOTA							1.00 70.21	6
MOTA	2659		1 ASP A 334	29.097	51.875	41.298	1.00 71.47	8
ATOM	2660	OD	2 ASP A 334	29.181	51.344	43.427	1.00 70.91	8
MOTA	2661	С	ASP A 334	26.660	52.041	44.627	1.00 65.65	6
ATOM	2662	ō	ASP A 334	26.797	50.996	43.990	1.00 63.91	
			•					8
ATOM	2663	Ŋ	PHE A 335	25.822	52.153	45.649	1.00 65.73	7
ATOM	2664	CA	PHE A 335	25.041	51.021	46.104	1.00 63.44	6
ATOM	2665	CB	PHE A 335	24.980	51.034	47.632	1.00 58.05	6
ATOM	2666	CG		24.039	50.028	48.195	1.00 53.82	6
	2667							
ATOM			1 PHE A 335	24.178	48.679	47.886	1.00 52.40	6
ATOM	2668		2 PHE A 335	22.978	50.429	48.989	1.00 51.33	6
ATOM	2669	CE.	1 PHE A 335	23.265	47.742	48.356	1.00 52.73	6
ATOM	2670	CE	2 PHE A 335	22.062	49.503	49.462	1.00 53.20	6
ATOM	2671	CZ	PHE A 335	22.204	48.151	49.144	1.00 51.76	6
MOTA	2672	C	PHE A 335	23.629	50.893	45.535	1.00 65.55	6
ATOM	2673	0	PHE A 335	23.230	49.810	45.097	1.00 67.33	8
ATOM	2674	N	GLU A 336	22.874	51.986	45.537	1.00 66.47	7
ATOM	2675	CA	GLU A 336	21.497	51.948	45.048	1.00 67.43	6
	2676	CB	GLU A 336	21.422	51.379	43.626	1.00 71.79	6
ATOM								
ATOM	2677	CG	GLU A 336	19.982	51.245	43.116	1.00 78.77	6
ATOM	2678	CD	GLU A 336	19.868	50.505	41.789	1.00 82.67	6
MOTA	2679	OE:	L GLU A 336	20.232	49.306	41.734	1.00 83.29	8
ATOM	2680	OE2	2 GLU A 336	19.410	51.126	40.801	1.00 84.26	8
	2681	C	GLU A 336	20.655	51.069			
MOTA						45.971	1.00 64.72	6
ATOM	2682	0	GLU A 336	20.686	49.840	45.876	1.00 59.84	8
ATOM	2683	N	GLU A 337	19.901.	51.710	46.858	1.00 64.47	7
ATOM	2684	CA	GLU A 337	19.045	51.003	47.805	1.00 65.83	6
ATOM	2685	CB	GLU A 337	18.398	52.003	48.759	1.00 64.20	6
	2686	CG	GLU A 337	17.753	51.370			
ATOM						49.964	1.00 64.26	6
ATOM	2687	CD	GLU A 337	18.774	0ز€.50	50.850	1.00 64.04	6
ATOM	2688	0E1		19.741	51.3.2	51.261	1.00 61.66	8
MOTA	2689	OE2	GLU A 337	18.608	49.483	51.132	1.00 63.64	8
ATOM	2690	С	GLU A 337	17.950	50.239	47.063	1.00 67.13	6
	2691			17.269	50.807			
MOTA		0	GLU A 337			46.205	1.00 68.27	8
ATOM	2692	N	PHE A 338	17.779	48.960	47.394	1.00 67.22	7
ATOM	2693	CA	PHE A 338	16.764	48.129	46.748	1.00 68.05	6
ATOM	2694	CB	PHE A 338	16.445	46.919	47.626	1.00 69.68	6
	2695	CG	PHE A 338	15.228	46.158	47.187	1.00 72.35	6
MOTA								9
ATOM	2696	CD1		15.122	45.674	45.888	1.00 72.37	6
MOTA	2697	CD2	PHE A 338	14.172	45.941	48.074	1.00 73.61	б
ATOM	2698	CE1	PHE A 338	13.980	44.984	45.478	1.00 73.39	6
	2699	CE2		13.024	45.250	47.672	1.00 73.26	6
ATOM								
ATOM	2700	CZ	PHE A 338	12.929	44.771	46.369	1.00 73.34	6
ATOM	2701	C	PHE A 338	15.481	48.902	46.434	1.00 68.45	6
ATOM	2702	0	PHE A 338	15.286	49.367	45.310	1.00 67.92	8
ATOM	2703	N	ASP A 339	14.606	49.026	47.426	1.00 68.98	7
				13.358				6
ATOM	2704	CA	ASP A 339		49.759	47.261	1.00 70.68	
ATOM	2705	CB	ASP A 339	12.596	49.758	48.588	1.00 71.06	6
ATOM	2706	CG	ASP A 339	11.381	50.678	48.581	1.00 72.05	6
						-		

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ATOM	2707	OD	1 ASP	A 339	11.548	51.893	48.320	1.00 72.29	0
ATOM	2708		2 ASP		10.262		48.858	1.00 71.03	8
ATOM	2709	c		A 339	13.715	51.183	46.853	1.00 71.03	8
ATOM	2710	ō		A 339	14.407	51.884	47.592	1.00 73.18	6
ATOM	2711	N		A 340	13.247	51.600	45.677	1.00 76.36	8
ATOM	2712	CA		A 340	13.518	52.943			7
ATOM	2713	CB		A 340	12.410		45.152	1.00 78.34	6
	2713	CG		A 340		53.385	44.189	1.00 77.55	6
MOTA	2715		1 ASP		12.462	52.655	42.864	1.00 78.90	6
ATOM					12.348	51.408	42.855	1.00 78.38	8
ATOM	2716		2 ASP		12.620	53.336	41.830	1.00 78.74	8
MOTA	2717	c		A 340 .	13.687	54.017	46.214	1.00 79.51	6
MOTA	2718	0		A 340	14.587	54.856	46.117	1.00 80.19	8
MOTA	2719	N		A 341	12.824	54.000	47.224	1.00 79.64	7
ATOM	2720	CA		A 341	12.922	54.998	48.271	1.00 80.05	6
MOTA	2721	CB		A 341	12.269	56.301	47.811	1.00 83.75	6
MOTA	2722	CG		A 341	12.411	57.442	48.806	1.00 89.02	6
ATOM	2723			A 341	11.756	58.724	48.328	1.00 91.52	6
ATOM	2724	OE:		A 341 .	10.515	58.738	48.175	1.00 93.33	. 8
MOTA	2725	OE2			12.484	59.716	48.102	1.00 92.65	8
ATOM	2726	C		A 341	12.317	54.578	49.597	1.00 77.98	6
MOTA	2727	0	GLU A		11.102	54.610	49.777	1.00 79.82	8
MOTA	2728	N	VAL A		13.179	54.181	50.523	1.00 74.49	7
MOTA	2729	CA	VAL A		12.745	53.793	51.859	1.00 71.55	6
ATOM	2730	CB	VAL A		13.224	52.383	52.245	1.00 72.40	6
ATOM	2731		. VAL ?		12.672	52.004	53.610	1.00 71.16	6
MOTA	2732		VAL A		12.797	51.391	51.207	1.00 74.35	6
ATOM	2733	С	VAL A		13.454	54.778	52.766	1.00 68.46	6
ATOM	2734	0	VAL ?		12.952	55.154	53.829	1.00 68.96	8
MOTA	2735	N	ASP A		14.636	55.184	52.311	1.00 61.61	7
ATOM-	2736	CA	ASP A		15.486	56.114	53.029	1.00 54.91	6
MOTA	2737	CB	ASP A		14.678	57.303	53.543	1.00 55.06	6
MOTA	2738	CG	ASP A	343	15.556	58.390	54.114	1.00 54.44	6
ATOM	2739		ASP A		15.002	59.351	54.694	1.00 56.20	8
ATOM	2740	QD2	ASP A		16.795	58.287	53.969	1.00 49.32	8
MOTA	2741	C	ASP A		16.152	55.401	54.198	1.00 50.85	6
ATOM	2742	0	ASP A	343	15.557	55.209	55.257	1.00 49.32	8
ATOM	2743	N	ARG A	344	17.396	55.004	53.980	1.00 47.84	7
MOTA	2744	CA	ARG A	344	18.195	54.321	54.981	1.00 45.34	6
ATOM	2745	CB	ARG A	344	18.883	53.099	54.358	1.00 45.00	6
ATOM	2746	CG	ARG A		17.950	51.969	53.974	1.00 38.03	6
ATOM	2747	CD	ARG A	344	17.185	51.531	55.188	1.00 35.83	6
ATOM	2748	NE	ARG A	. 344	16.278	50.439	54.885	1.00 39.20	7
MOTA	2749	CZ	ARG A		15.350	49.993	55.724	1.00 39.89	6
ATOM	2750	NH1	ARG A	344	15.217	50.561	56.917	1.00 40.17	7
ATOM	2751	NH2	ARG A	344	14.566	48.976	55.375	1.00 40.75	7
ATC 1	2752	С	ARG A	344	19.250	55.278	55.515	1.00 44.72	6
ATC:M	2753	0	ARG A	344	20.170	54.869	56.223	1.00 46.97	8
ATOM	2754	N	SER A		19.113	56.552	55.157	1.00 45.81	7
ATOM	2755	CA	SER A	345	20.045	57.596	55.577	1.00 43.66	6
ATOM	2756	CB	SER A	345	19.538	58.960	55.115	1.00 43.44	6
ATOM	2757	OG	SER A	345	18.292	59.260	55.722	1.00 45.62	8
ATOM	2758	С	·SER · A	345	20.258	57.627	57.089	1.00 42.79	6
MOTA	2759	0	SER A	345	21.364	57.902	57.5 52	1.00 42.62	8
ATOM	2760	N	TYR A	346	19.200	57.354	57.851	1.00 40.55	7
ATOM	2761	CA	TYR A	346	19.280	57.352	59.308	1.00 41.05	6
ATOM	2762	CB	TYR A	346	17.971	56.819	59.905	1.00 41.74	6
ATOM	2763	CG	TYR A		17.668	55.355	59.630	1.00 43.47	6
ATOM	2764	CD1			18.331	54.333	60.328	1.00 44.45	6
ATOM	2765		TYR A		18.044	52.983	60.088	1.00 41.02	6
ATOM	2766	CD2	TYR A		16.710	54.988	58.682	1.00 42.08	6
ATOM ATOM	2767	CE2	TYR A		16.416	53.644	58.434	1.00 40.59	6
ATOM . ATOM	2768	CZ	TYR A		17.086		59.139	1.00 41.66	5
	2769	ОН	TYR A	_	16.806	51.324	58.884	1.00 39.60	8
ATCM	2770	C	TYR A		20.466	.56.517	59.796	1.00 42.92	6
ATCM	2771	ō	TYR A		21.101	56.844	60.799	1.00 42.65	8
ATCM		N	MET A		20.757		59.067	1.00 44.59	7
ATCM	2772	44			20.737	55.443	-	7.00 44.73	'

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Figure 17-43

									•
ATOM	2773	CA	MET A 347		21.859	54.546	59.388	1.00 45,89	6
MOTA	2774				21.950		58.353	1.00 45.51	6
MOTA	. 2775	CG	MET A 347		20.727	52.567	58.244	1.00 45.01	6
ATOM	2776	SE	MET A 347		21.062	51.258	57.066	1.00 47.13	16
							-		
ATOM	2777				21.545	52.237	55.676	1.00 44.55	6
ATOM -	2778	С	MET A 347		23.188	55.286	59.404	1.00 48.47	6
ATOM	2779	0	MET A 347		24.129	54.888	60.098	1.00 49.49	8
MOTA	2780		LEU A 348		23.259	56.351	58.610	1.00 49.03	7
MOTA	2781	. CA	LEU A 348		24.458	57.178	58.499	1.00 48.86	6
-	2782	CB	LEU A 348		24.355	58.082	57.269	1.00 45.51	6
ATOM									
ATOM	2783				24.280	57.424	55.895	1.00 44.47	6
MOTA	2784	CD	1 LEU A 348		23.908	58.476	54.859	1.00 43.62	6
ATOM	2785	CD	2 LEU A 348	-	25.618	56.757	55.565	1.00 42.53	6
			LEU A 348						
MOTA	2786				24.644	58.049	59.738	1.00 49.33	6
ATOM	2787	0	LEU A 348		25.765	58.369	60.123	1.00 49.78	8
MOTA	2788	N	GLU A 349		23.537	58.428	60.358	1.00 48.34	7 -
	2789				23.591	59.279	61.533		6
MOTA								1.00 49.24	
ATOM	2790	CB	GLU A 349		22.198	59.848	61.811	1.00 48.36	6
ATOM	2791	CG	GLU A 349		21.628	60.584	60.607	1.00 45.52	6
	2792		GLU A 349		22.598	61.619	60.065	1.00 42.94	6
ATOM									
ATOM	2793		1 GLU A 349		22.934	62.560	60.812	1.00 40.82	8
MOTA	2794	OE.	2 GLU A 349		23.028	61.483	58.900	1.00 38.30	8
ATOM	2795	С	GLU A 349		24.119	58.531	62.745	1.00 48.32	6
	2796	ō	GLU A 349		25.226	58.783	63.219	1.00 47.87	8
ATOM									
MOTA	2797	N	THR A 350		23.325	57.602	63.248	1.00 48.97	7
MOTA	2798	CA	THR A 350		23.744	56.832	64.398	1.00 50.70	6
ATOM	2799	CB	THR A 350		22.558	56.596	65.342	1.00 51.02	6
	2800	OG:	•		22.071	57.865	65.803	1.00 49.11	8
MOTA									
ATOM	2801	CG:			22.983	55.763	66.537	1.00 51.58	6
MOTA	2802	C	THR A 350		24.361	55.507	63.954	1.00 49.56	б
MOTA	2803	0	THR A 350		23.979	54.947	62.923	1.00 50.55	8
	2804	N	LEU A 351		25.333	55.028	64.725	1.00 46.88	7
ATOM									
ATOM	2805	CA	LEU A 351		26.018	53.781		1.00 45.35	6
MOTA	2806	CB	LEU A 351		27.342	53.726	65.185	1.00 47.05	6
MOTA	2807	CG	LEU A 351		28.257	52.502	65.072	1.00 49.54	6
ATOM	2808	CDI			29.575	52.777	65.766	1.00 51.50	6
MOTA	2809		LEU A 351		27.603	51.302	65.692	1.00 48.35	6
ATOM	2810	С	LEU A 351		25.145	52.584	64.772	1.00 44.79	6
MOTA	2811	0	LEU A 351		25.131	51.578	64.061	1.00 41.45	8
ATOM .	2812	N	LYS A 352		24.420	52.711	65.880	1.00 45.27	7
			LYS A 352				66.375	1.00 44.62	6
ATOM	2813	CA			23.531	51.662			
ATOM	2814	CB	LYS A 352		23.764	51.464	67.873	1.00 42.23	6
MOTA	2815	CG	LYS A 352		25.197	51.075	68 .187	1.00 44.94	6
ATOM	2816	CD	LYS A 352		25.572	51.262	69 .650	1.00 46.80	6
	2817		LYS A 352						6
ATOM		CE			24.765	50.389	70.581	1.00 45.79	
ATOM	2818	NZ	LYS A 352		25.236	50.586	71 .97 5	1.00 47.31	7
ATOM	2819	С	LYS A 352		22.096	52.087	66.11 6	1.00 45.12	6
MOTA	2820	0	LYS A 352		21.837	53.236	65.756	1.00 47.07	8
	2821	Ŋ	ASP A 353		21.162	51.161	66.285	1.00 44.62	7 .
MOTA									
ATOM	2822	CA	ASP A 353		19.761	51.474	66.060	1.00 46.43	6
MOTA	2823	CB	ASP A 353		19.302	50.943	64.692	1.00 49.38	6
	.2824.	CG	ASP A 353		19.813	49.546	64.396	.1.00 51.52	6
MOTA									
ATOM	2825		ASP A 353		21.028	49.396	64.158	1.00 55.36	8
ATOM	2826	OD2	ASP A 353		19.005	48.596	64.398	1.00 52.35	8
ATOM	2827	С	ASP A 353		18.841	50.968	67.165	1.00 45.90	6
	2828	ō	ASP A 353		19.152	50.001	67.854	1.00 45.98	8
ATOM									7
ATOM	2829	N	PRO A 354		17.687	51.629	67.348	1.00 45.86	7
ATOM	2830	CD	PRO A 354		17.162	52.775	66.587	1.00 45.36	6
	2831	CA	PRO A 354		16.723	51.243	68.378	1.00 45.52	6
MOTA								1.00 44.77	6
ATOM	2832	CB	PRO A 354		15.585	52.245	68.159		
MOTA	2833	CG	PRO A 354		15.681	52.513	66.664	1.00 45.06	6
ATOM	2834	С	PRO A 354		16.277	49.804	68. 188	1.00 44.13	6
	2835	Ö	PRO A 354		16.352	49.271	67.078	1.00 42.90	8
ATOM			TRP A 355		15.821		69.267	1.00 42.77	7
ATCM	2836	N				49.174			
ATOM	2837	CA	TRP A 355		15.358	47.801	69. 168	1.00 43.35	6
ATOM	2838	CB	TRP A 355		14.982	47.225	70.539	1.00 47.11	6
			_						

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ATO		16.168 46.752 71.322 1.00 52 43 6	
ATO		16 574 45 391 71 510 1 00 52.43 6	
ATO		17.789 45.416 72.238 1 00 54 97 6	
ATO		16 031 44 153 71 155 7 75	
ATOI	32 333	17.125 47.526 71.916 1.00 54.39 6	
OTA		18.103 46.731 72.468 1.00 57.31 7	
ATON		18.469 44.249 72.602 1.00 54 97 6	
1OTA 1OTA		16.706 42.995 71.518 1.00 55.77 6	
ATOM		17.913 43.052 72.234 1.00 54.84 6	
MOTA	555	14.177 47.690 68.230 1.00 41.94 6	
ATOM		13.508 48.677 67.915 1.00 41.39 8	
ATOM		13.942 46.471 67.775 1.00 38.60 7	
ATOM		12.855 46.185 66.866 1.00 36.55 6	
АТОМ		13.413 46.044 65.451 1.00 35.06 6	
ATOM		14.120 47.308 64.976 1.00 32.47 6	
ATOM		14.969 47.082 63.733 1.00 29.54 6	
ATOM	12.0 11. 330	1.00 20.31	
ATOM	2857 NH1 APC 3 356	16.514 48.403 62.335 1.00 30.50 6 16.916 47.305 61.702 1.00 33.53	
ATOM		1.00 33.32.	
ATOM	2859 C ARG A 356	10 30.10	
ATOM	2860 O ARG A 356	12 447 43 931 66 742 1 00 56.01 6	
MOTA	2861 N GLY A 357	11 507 44 040 60 100 30.36 8	
ATOM	2862 CA GLY A 357	1.00 30.04	
ATOM	2863 C GLY A 357	9 514 43 506 60 051	
MOTA	2864 O GLY A 357	8.943 44.196 67.943 1 00 36 77 0	
ATOM	2865 N GLY A 358	8.892 42.772 69.687 1.00 36.04 7	
ATOM	2866 CA GLY A 358	7.466 42.506 69.593 1.00 32.26 6	
ATOM ATOM	2867 C GLY A 358 2868 O GLY A 358	7.106 41.263 70.385 1.00 29.85 6	
ATOM		7.832 40.839 71.288 1.00 28.86 8	
ATOM		5.975 40.667 70.055 1.00 30.88 7	
ATOM	2870 CA GLU A 359 2871 CB GLU A 359	3.550 39.455 70.743 1.00 32.58 6	
ATOM	2872 CG GLU A 359	4.034 39.289 70.604 1.00 38.60 6	
ATOM	2873 CD GLU A 359	3.230 40.435 71.222 1.00 47.44 6 1.957 40.762 70 445 1.00 50 03	
ATOM	2874 OE1 GLU A 359	10.330 7.00 30.33 6	
ATOM	2875 OE2 GLU A 359		
ATOM	2876 C GLU A 359	6 250 30 275 70 201	
MOTA	2877 O GLU A 359	6 790 30 303 60 000 1 00 20.29 6	
ATOM	2878 N VAL A 360	6.263 37.147 70.772 1.00 27.88 8	
ATOM	2879 CA VAL A 360	6.859 35.957 70.193 1.00 25.86 6	
ATOM	2880 CB VAL A 360	7.6/3 35.168 71.237 1.00 22.02 6	
ATOM ATOM	2881 CG1 VAL A 360 2882 CG2 VAL A 360	8.155 33.849 70.641 1.00 19.45 6	
ATOM		8.850 36.009 71.698 1.00 17 88 6	
ATOM	2883 C VAL A 360 2884 C VAL A 360	5.703 35.099 69.670 1.00 28.04 6	
ATOM	2885 N ARG A 361	4.842 34.655 70.440 1.00 27.34 8	
ATOM -	2886 C ARG A 361	5.663 34.898 68.358 1.00 27.70 7 4.612 34.091 67.765 1.00 32.95	
MOTA	2887 CB ARG A 361	2,00 32,05 6	
MOTA	2888 CG ARG A 361	1.00 32.30	
MOTA	2889 CD ARG A 361	4 546 35 553 64 301 1.00 30.81 8	
MOTA	2890 NE ARG A 361		
ATOM	2891 CZ ARG A 361	6 514 35 000	
ATOM	2892 NH1 ARG A 361	5 749 36 937 51 535	
ATOM	2893 NH2 ARG A 361	7 922 36 145 62 642	
ATOM	2894 C ARG A 361	4 689 33 638 60 333	
MOTA	2895 O ARG A 361	5.768 32.097 68.471 1.00 37.08 8	
MOTA	2896 N LYS A 362	3.526 32.017 68.347 1.00 37.80 7	
MOTA	2897 CA LYS A 362	3.436 30.626 68.757 1.00 39.91 6	
MOTA	2898 CB LYS A 362	1.982 30.152 68.648 1.00 43.43 6	
MOTA	2899 CG LYS A 362	1.014 30.803 69.640 1.00 45.11 6	
ATOM TOM	2900 CD LYS A 362 2901 CE LYS A 362	1.117 32.346 69.673 1.00 49.43 6	
MOTA MOTA	2901 CE LYS A 362 2902 NZ LYS A 362	0.813 33.022 68.327 1.00 45.02 6	
TOM	2903 C LYS A 362	0.962 34.509 68.422 1.00 41.08 7	
ATOM	2904 O LYS A 362	4.320 29.809 67.831 1.00 39.25 6	
		4.953 28.835 68.248 1.00 35.45 8	

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ATO	M 2905 N GLU A 363	4.358 30.229 66.568 1 00 41 17
ATO		5 1.13
ATOI		7 00 43.96
ATO	M 2908 CG GLU A 363	3 300
ATO		1 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
ATON		1.00 34.11
OTA		3 640 33 80 8
ATON		, tan a
ATON		7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7
ATOM		7 10 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2
ATOM		2.00 37.27
ATOM		2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2
· ATOM	2917 CG1 VAL A 364	9.038 31.696 67.469 1.00 36.88 6
ATOM		10.444 31.599 68.043 1.00 37.77 6
ATOM		9.018 32.628 66.252 1.00 34.78 6
ATOM		8.650 29.361 68.268 1.00 33.01 6
ATOM		9.622 28.614 68.379 1.00 31.55 8
ATOM		7.664 29.409 69.165 1.00 32.61 7
ATOM		7.674 28.567 70.362 1.00 30.96 6
ATOM		6.598 29.010 71.358 1.00 30.13 6
ATOM		0.826 30.409 71.899 1.00 36.02 6
ATOM	500	3.837 30.781 72.995 1.00 38.94 6
ATOM		6.120 32.187 73.509 1.00 41.58 6
ATOM		3.191 32.611 74.585 1.00 44.29 7
ATOM		7.452 27.114 70.007 1.00 30.32 6
ATOM		8.195 26.237 70.442 1.00 31 32 0
ATOM		0.42/ 20.863 69.209 1.00 29 85 7
ATOM		0.115 25.509 68.807 1.00 32.07 6
ATOM		4.948 25.522 67.818 1 00 35 00 6
		3./11 26.206 68.381 1 00 39 99 6
ATOM		3.124 25.692 69.359 1.00 40 16 9
ATOM	2935 OD2 ASP A 366	3.326 27.266 67.848 1.00 43.21 R
ATOM	2936 C ASP A 366	7.343 24.866 68.161 1 00 33 03
ATOM	2937 O ASP A 366	7.753 23.763 68.540 1.00 32 64 8
MOTA	2938 N THR A 367	7.932 25.565 67.193 1 00 31 90 7
ATOM	2939 CA THR A 367	9.088 25.045 66.490 1.00 31.17 6
ATOM	2940 CB THR A 367	9.712 26.070 65.572 1.00.31.55 6
ATOM	2941 OG1 THR A 367	8.707 26.620 64.714 1.00 34.37 8
ATOM	2942 CG2 THR A 367	10.780 25.404 64.723 1.00 33.55 6
MOTA	2943 C THR A 367	10.146 24.633 67.472 1.00 33.09 6
ATOM	.2944 O THR A 367	10.586 23.476 67.485 1.00 38.62 8
ATOM	2945 N LEU A 368	10.570 25.579 68.298 1.00 31.85 7
ATOM	2946 CA LEU A 368	11.582 25.264 69.288 1.00 32.87 6
ATOM	2947 CB LEU A 368	11.848 26.478 70.179 1.00 27.73 6
ATOM	2948 CG LEU A 368	12.887 27.449 69.588 1.00 29.05 6
MOTA	2949 CD1 LEU A 368	14.260 26.777 69.541 1.00 23.05 6
MOTA	2950 CD2 LEU A 368	12.473 27.896 68.193 1.00 26.53 6
ATOM	2951 C LEU A 368	11.157 24.053 70.107 1.00 35.16 6
ATOM	2952 O LEU A 368	11.910 23.077 70.217 1.00 35.18 8
ATOM	2953 N GLU A 369	9.942 24.101 70.649 1.00 37.56 7
ATOM	2954 CA GLU A 369	9.431 22.993 71.442 1.00 40.23 6
MOTA	2955 CB GLU A 369	7.956 23.216 71.770 1.00 42.07 6
MOTA	2956 CG GLU A 369	7.722 24.460 72.617 1.00 48.51 6
ATOM	2957 CD GLU A 369	6 791 74 616 77 965
ATOM	2958 OE1 GLU A 369	5 777 23 724 72 720 4 24
ATOM	2959 OE2 GLU A 369	5 652 25 626 70 712
MOTA	2960 C GLU A 369	9 633 23 672 70 70
ATOM	2961 O GLU A 369	10 087 20 604 71 206
ATOM	2962 N LYS A 370	0 300 0: 650
MOTA	2963 CA LYS A 370	0 407 00 440 00 100
MOTA	2964 CB LYS A 370	9 144 20 654 67 66 1 00 10.20
MOTA	2965 CG LYS A 370	7 675 20 507 56 054 1 00 11
ATOM	2966 CD LYS A 370	7 495 20 250 65 262 1 20 40 25
ATOM	2967 CE LYS A 370	6 052 20 027 05 015
ATOM	2968 NZ LYS A 370	E 890 10 CTO CT TTO 100 ST. 20
ATOM	2969 C LYS A 370	
ATOM	2970 O LYS A 370	10.948 20.034 68.730 1.00 37.85 6
	DIO O LIS A 3/0	11.261 18.930 69.156 1.00 37.95 8
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	2077							
ATOM	2971		A 371	11.827		68.325	1.00 37.78	7
ATOM	2972	CA ALA	A 371	13.264	20.704	68.340	1.00 36.39	6
ATOM	2973	CB ALA	A 371	14.007	22.030	68.200	1.00 37.73	6
ATOM	2974		A 371	13.719	19.972	69.603	1.00 35.42	
	2975		_					6
ATOM			A 371	14.424	18.964	69.525	1.00 32.06	8
MOTA	2976		A 372	13.317	20.478	70.766	1.00 33.96	7
ATOM	2977	CA ALA	A 372	13.695	19.848	72.024	1.00 32.22	6
ATOM	2978	CB ALA	A 372	12.946	20.486	73.165	1.00 28.27	6
MOTA	2979		A · 372	13.372	18.362	71.953		
							1.00 31.75	6
ATOM	2980		A 372	14.183	17.517	72.338	1.00 31.56	8
ATOM	2981		¥ 373	- 12.187	18.059	71.432	1.00 32.72	7 .
MOTA	2982	CA ALA A	373	11.710	16.684	71.305	1.00 32.32	6
MOTA	2983	CB ALA	373	10.206	16.689	71.103	1.00 30.18	6
ATOM	2984	C ALA		12.385	15.921	70.172	1.00 33.13	
	2985	OT1 ALA						6
MOTA				13.078	14.926	70.468	1.00 35.87	8
MOTA	2986	OT2 ALA:		12.218	16.320	69.003	1.00 34.11	8
ATOM	2987	ZN ZN 2	2 951	22.693	34.497	53.990	1.00 36.45	6
ATOM	2988	OH2 WAT S	3 1	35.654	44.211	49.416	1.00 9.27	8
ATOM	2989	OH2 WAT S		24.480	33.130	53.069	1.00 21.27	8
ATOM	2990	OH2 WAT S		22.124	30.277	59.314		
		•					1.00 14.69	8
ATOM	2991	OH2 WAT S		13.839	20.611	75.741	1.00 27.94	8
ATOM	2992	OH2 WAT S		34.033	41.903	46.522	1.00 44.54	8
MOTA	2993	OH2 WAT S	6	15.039	42.130	55.781	1.00 23.79	8
ATOM	2994	OH2 WAT S	7	32.737	41.397	75.900	1.00 15.80	8
ATOM	2995	OH2 WAT S	8	11.367	22.606	58.814	1.00 23.37	8
ATOM	2996	OH2 WAT S	9	13.909	18.160	65.105	1.00 29.93	8
ATOM	2997	OH2 WAT S	-	29.655	56.108	58.029	1.00 50.54	
ATOM	2998	OH2 WAT S						8
				45.405	17.964	51.885	1.00 9.28	8
ATOM	2999	OH2 WAT S		21.870	35.873	34.515	1.00 32.78	8
ATOM	3000	OH2 WAT S		43.504	35.670	33.779	1.00 28.85	8
ATOM	3001	OH2 WAT S		2.054	37.997	68.430	1.00 40.53	8
MOTA	3002	OH2 WAT S	15	49.730	28.024	55.966	1.00 21.42	.8
ATOM	3003	OH2 WAT S	16	47.503	32.289	34.336	1.00 26.13	8
ATOM	3004	OH2 WAT S	17	6.101	26.102	64.434	1.00 21.69	8
ATOM	3005	OH2 WAT S	_	10.761	46.748	45.836	1.00 15.79	8
ATOM	3006	OH2 WAT S		9.146	16.861			
						61.441	1.00 16.68	8
ATOM	3007	OH2 WAT S		5.684	34.080	76.599	1.00 37.53	8
ATOM	3008	OH2 WAT S	21	14.896	33.163	49.117	1.00 34.17	8
MOTA	3009	OH2 WAT S	22	43.346	40.839	36.825	1.00 35.64	8
ATOM	3010	OH2 WAT S	23	0.516	27.705	69.174	1.00 21.02	8
ATOM	3011	OH2 WAT S	24	41.270	25.444	29.717	1.00 29.80	8
MOTA	3012	OH2 WAT S	25	17.818	29.142	54.584	1.00 27.92	8
ATOM	3013	OH2 WAT S	26	21.512	60.572	56.912	1.00 16.77	8
ATOM	3014	OH2 WAT S	27	21,211		48.347		
							1.00 23.93	8
ATOM	3015	OH2 WAT S	28	47.805	24.638	56.619	1.00 23.73	8
MOTA	3016	OH2 WAT S	25	44.624	50.302	58.154	1.00 16.79	8
ATOM	3017	OH2 WAT S	3€	31.096	16.437	51.311	1.00 26.61	8
ATOM	3018	OH2 WAT S	31	39.837	38.833	55.145	1.00 32.28	8
MOTA	3019	OH2 WAT S	· 32	11.660	43.601	63.704	1.00 22.94	8
MOTA	3020	OH2 WAT S	33	49.899	23.474	53.058	1.00 26.85	8
	3021	OH2 WAT S			17.734		1.00 21.18	8
	3022	OH2 WAT S	35	26.926	15.913	62.444		
ATOM							1.00 27.01	8
ATOM	3023	OH2 WAT S	36	8.893	28.686	63.905	1.00 27.68	8
ATOM	3024	OH2 WAT S	37	23.381	26.634 ⁻	43.532	1.00 24.42	8
ATOM	3025	OH2 WAT S	38	48.484	27.990	65.270	1.00 34.86	8
ATOM	3026	OH2 WAT S	39	43.382	28.410	74.379	1.00 25.68	8
ATOM	3027	OH2 WAT S	40	42.904	18.967	70.272	1.00 29.45	8
ATOM	3028	OH2 WAT S	41	20.521	53.828	50.298	1.00 22.35	8
	3029	OH2 WAT S	42	13.310		48.404		
ATOM					38.921		1.00 23.32	8
MOTA	3030	OH2 WAT S	43	9.787	46.265	60.012	1.00 33.51	8
ATOM	3031	OH2 WAT S	44	36.089	30.416	51.377	1.00 47.75	8
ATOM	3032	OH2 WAT S	45	14.831	48.131	42.151	1.00 50.96	8
ATOM	3033	OH2 WAT S	46	54.162	48.194	60.971	1.00 22.66	8
ATOM	3034	OH2 WAT S	47	38.943	61.290	63.509	1.00 33.73	8
ATOM	3035	OH2 WAT S	48	29.980	18.112	33.130	1:00 35.80	8
•	3036	OH2 WAT S	49	31.879	50.673	44.528	1.00 24.39	8
ATOM	2020				55.575			~

ATOM	3037	OH2 V	VAT S	5 50		39.86	3 14.6	29 64.30	7 1 0	0 24.19	` .
ATOM	3038					26.11				0 24.15	
ATOM				52		48.07	0 41.5			36.38	8
ATOM		OH2 W				50.80	2 29.6		5 1.0	31.04	8
ATOM		OH2 W		_		49.54	0 35.5	32 71.58	5 1.0	20.96	8
MOTA		OH2 W				6.88			1 1.00	17.49	8
ATOM		OH2 W				25.69		37.67	4 1.00	51.51	. 8
ATOM		OH2 W		-		45.49			3 1.00	37.34	8
MOTA		OH2 W				44.66			2 1.00	44.52	8
ATOM		OH2 W		_		21.91			3 1.00	26.96	. 8
ATOM		OH2 W				27.29			1.00	27.74	8
MOTA MOTA		OH2 W			-	19.809				46.14	8
ATOM	-	OH2 W		62 63		30.843				42.23	8
MOTA	3051	OH2 W		64		19.059 47.929			1.00	28.99	8.
MOTA	3052	OH2 W		65		32.500				34.93	8
ATOM	3053	OH2 W		66		27.245				35.33	8
ATOM	3054	OH2 W		67		5.176			1.00	34.19	8
ATOM	3055	OH2 W		68	•	41.159			1.00	41.89	8
ATOM	3056	OH2 W		69		12.869				27.31	8
ATOM	3057	OH2 W	_	70		17.499				31.30 24.91	8
ATOM	3058	OH2 WA	AT S	71		27.152				18.76	8
MOTA	3059	OH2 WA	T S	72		25.213				61.35	8 8
ATOM	3060	OH2 WA		73		17.671			1.00	37.63	8
ATOM	3061	OH2 WA		74		23.765	60.84	6 66.579	1.00	21.81	8
ATOM	3062	CH2 WA	_	75		35.535		0 70.698		34.04	. 8
ATOM	3063	OH2 WA		76		26.280			1.00	32.20	. 8
MOTA MOTA	3064 3065	OH2 WA		77		18.451	25.55			28.55	8
ATOM	3066	OH2 WA		78 70		10.446	61.27			44.74	8
ATOM	3067	OH2 WA	_	79 80		13.256	24.05			35.45	8
ATOM	3068	OH2 WA		81		23.571 29.891	13.292 18.071		1.00	49.49	8
ATOM	3069	OH2 WA		82		12.886	42.723			22.84	8
ATOM	3070	OH2 WA		83 .		41.348	15.471			35.31	8
ATOM	3071	OH2 WA		84		13.406	44.647		1.00	47.24 49.67	8
MOTA	3072	OH2 WA		85		30.444	35.217			38.15	8
ATOM	3073	OH2 WA	T S	86		5.217	40.817			19.51	8 8
ATOM	3074	CH2 WA	T S	87		8.891	21.532			30.72	8
MOTA	3075	OH2 WA!		88		41.816	25.022			22.92	8
ATOM		OH2 WA		89		50.621	36.644	60.248		29.29	8
ATOM		OH2 WAT		90		26.008	34.532			45.42	8
ATOM		OH2 WAT		91		8.131	39.168			31.50	8
MOTA MOTA		OH2 WAT		92		16.591	58.091			34.73	8
MOTA		OH2 WAT		93 94		34.773	54.065			36.05	8
MOTA	_	OH2 WAT		95		42.105 29.684	31.720			35.49	8
ATOM		OH2 WAT		96	٠.	26.411	52.077 37.426			35.17	8
ATOM		OH2 WAT		97	•	41.183	52.989	38.934 62.927		41.68	8
ATOM		OH2 WAT		98		21.167	6.202	63.102	1.00		8
ATOM		OH2 WAT		99		25.060	18.985	36.669	1.00		8
ATOM	3087 (OH2 WAT	' S 1	00		37.304	39.027	73.722	1.00	: :	8 8
MOTA	3088	TAW SHC	S 1	01		15.911	54.635	39.343	1.00		8
MOTA	3089.	CH2 WAT	S 1	02 -		48.730	25.803	59.572	1.00		8
MOTA		DH2 WAT	S 1	03		24.029	42.997	74.111	1.00		8
MOTA		H2 WAT				42.477	21.773	46.986	1.00		8
ATOM		H2 WAT				29.984	22.945	31.397	1.00		8
MOTA	3093	H2 WAT	S 1	06		40.850	36.936	31.885	1.00		8
MOTA		H2 WAT				9.750	32.487	48.823	1.00		8
MOTA	3095 0	H2 WAT	S 10	80		7.618	30.171	58.896	1.00	40.03	8
ATOM	3096 0	H2 WAT	S 10	9		17.603	13.771	59.767	1.00		8
ATOM		H2 WAT				22.590	8.744	67.501	1.00		8
ATOM		H2 WAT				21.034	29.771	76.056	1.00		8
ATOM TOW		H2 WAT				24.791 40.750	14.674	50.081	1.00		8
ATOM		H2 WAT				40.750 7.708	47.494	54.056	1.00 4		8
MOTA MOTA		H2 WAT		_		32.375	42.479	58.027	1.00 3		8
J. Cur	-104 U	· · ·	3				49.136	77.566	1.00 2	27.53	8
						· ·					_

ATOM	3103	OH2 WAT S 116	5.596	17.009	64.551	1.00 39.15	8
ATOM	3104	OH2 WAT S 117	20.194	5.0.998	70.563	1.00 19.73	8
ATOM	3105	OH2 WAT S 118	23.853	64.927	64.164	1.00 27.16	8
ATOM	3106	OH2 WAT S 119	9.277	43.601	46.279	1.00 32.31	8
ATOM	3107	OH2 WAT S 120	15.613	24.398	46.723	1.00 55.20	8
ATOM	3108	OH2 WAT S 121	33.110	16.122	54.229	1.00 35.91	8
ATOM	3109	OH2 WAT S 122	26.772	34.085	33.852	1.00 37.49	8
ATOM	3110	OH2 WAT S 123	28.654	37.783	75.829	1.00 47.30	8
ATOM	3111	OH2 WAT 5 124	49.180	22.653	59.678	1.00 37.33	8
MOTA	3112	OH2 WAT S 125	20.561	27.788	65.975	1.00 67.86	8
ATOM	3113	OH2 WAT S 126	34.251	13.344	57.366	1.00 36.18	8
MOTA	3114	OH2 WAT S 127	49.215	36.854	48.117	1.00 33.63	8
ATOM	3115	OH2 WAT S 128	45.826	19.588	41.601	1.00 44.07	8
ATOM	.3116	OH2 WAT S 129	18.693	56.382	64.014	1.00 47.77	8
ATOM	3117	OH2 WAT S 130	44.181	24.202	36.963	1.00 32.70	8
ATOM	3118	OH2 WAT S 131	19.160	51.901	38.133	1.00 54.07	8
ATOM	3119	OH2 WAT S 132	16.904	36.558	48.679 34.353	1.00 42.21 1.00 56.33	. 8 . 8
MOTA	3120	OH2 WAT S 133	. 46.851 3.925	26.029 41.533	68.647	1.00 45.99	. 8
ATOM	3121	OH2 WAT S 134 OH2 WAT S 135	44.590	38.382	78.167	1.00 44.50	8
MOTA	3122 3123	OH2 WAT S 135	6.384	19.317	71.166	1.00 28.17	8
MOTA	3123	OH2 WAT S 130	17.982	39.823	66.487	1.00 49.31	8
ATOM ATOM	3125	OH2 WAT S 138	8.317	22.286	61.863	1.00 43.42	8
ATOM	3126	OH2 WAT S 139	29.248	14.196	55.622	1.00 35.55	8
MOTA	3127	OH2 WAT S 140	30.377	33.180	80.320	1.00 43.94	8
ATOM	3128	OH2 WAT S 141	41.842	32.906	27.392	1.00 24.82	8
MOTA	3129	OH2 WAT S 142	33.971	3.859	64.002	1.00 41.93	8
ATOM	3130	OH2 WAT S 143	27.314	8.087	70.916	1.00 49.03	8
ATOM	3131	OH2 WAT S 144	4.310	39.006	64.550	1.00 32.70	8
ATOM	3132	OH2 WAT S 145	2.940	19.950	63.265	1.00 33.24	8
ATOM	3133	OH2 WAT S 146	24.134	47.625	60.121	1.00 44.24	8
ATOM	3134	OH2 WAT S 147	25.035	53.746	42.337	1.00 47.82	8
MOTA	3135	OH2 WAT S 148	32.767	38.897.		1.00 21.86	8
MOTA	3136	OH2 WAT S 149	37.145	57.288	47.392	1.00 36.13 1.00 38.04	8 8
MOTA	3137	OH2 WAT S 150	25.171 24.054	18.011 43.182	32.273 55.583	1.00 38.04	8
MOTA	3138	OH2 WAT S 151 OH2 WAT S 152	27.686	64.936	52.937	1.00 60.62	8
ATOM	3139 3140	OH2 WAT S 152	24.084	39.543	76.589	1.00 22.62	8
ATOM	3141	OH2 WAT S 154	42.110	10.159	68.662	1.00 46.98	8
MOTA MOTA	3142	OH2 WAT S 155	9.675	22.905	75.335	1.00 26.45	8
ATOM	3143	OH2 WAT S 156	4.506	34.799	52.857	1.00 33.84	8
ATOM	3144	OH2 WAT S 157	32.583	35.051	76.446	1.00 36.27	8
ATOM	3145	OH2 WAT S 158	40.341	58.311	60.390	1.00 54.69	8
ATOM	3146	OH2 WAT S 159	29.473	58.378	71.881	1.00 28.59	8
ATOM	3147	OH2 WAT S 160	11.829	60.543	56.138	1.00 37.67	8
ATOM	3148	OH2 WAT S 161	24.247	48.010	67.935	1.00 56.62	8
ATOM	3149	OH2 WAT S 162	12.85	33.929	77.503	1.00 29.88	8
MOTA	3150	OH2 WAT S 163	9.49`	26.168	59.687	1.00 15.42	8
ATOM	3151	OH2 WAT 5 164	27.424	16.480	38.895	1.00 36.86	8
ATOM	3152	OH2 WAT S 165	8.512	56.634	49.614	1.00 30.08	8 8
MOTA	3153	OH2 WAT S 166	30.721	13.394 38.223	73.903	1.00 39.47	8
ATOM	3154	OH2 WAT S 167	49.594 41.994	48.023	74.119	1.00 38.12	8
ATOM	3155	OH2 WAT S 168 OH2 WAT S 169	42.092	39.503	33.116	1.00 24.47	8
ATOM	3156	OH2 WAT S 109 OH2 WAT S 170	34.547	12.749	38.054	1.00 38.65	8
MOTA	3157	OH2 WAT S 170	15.377	60.862	50.791	1.00 32.82	8
ATOM	3158 3159	OH2 WAT S 171	31.854	42.110	62.950	1.00 42.43	8
ATOM	3159	OH2 WAT S 172	48.743	44.073	57.626	1.00 34.04	8
ATOM	3161	OH2 WAT S 174	8.723	50.038	42.232	1.00 32.87	8
MOTA MOTA	3162	OH2 WAT S 175	14.257	18.280	53.455	1.00 40.51	8
ATOM	31.63	OH2 WAT S 176	31.917	37.509	53.943	1.00 40.43	8
ATOM	3164	OH2 WAT S 177	23.921	47.029	70.642	1.00 47.97	8
MOTA	3165	OH2 WAT S 178	27.974	47.778	69.949	1.00 62.12	8
ATOM .		OH2 WAT S 179	7.850	25.093	51.345	1.00 50.13	8
ATOM	3167	OH2 WAT S 180	22.080	48.840	66.463	1.00 53.81	8
ATOM	3168	OH2 WAT S 181	34.780	48.220	77.419	1.00 30.86	8

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Figure 17-49

MOTA	3169	OH2 WAT S 182	43,893.35	.526 52.018	1.00 47.14	8
ATOM	3170	OH2 WAT S 183	29.166 21	.424 28.950	1.00 45.08	8
ATOM	3171	OH2 WAT S 184	51.175 51	.545 62.599	1.00 33.88	8
ATOM	3172	OH2 WAT S 185	18.520 46	.208 42.323	1.00 50.85	8
MOTA	3173	OH2 WAT S 186	44.774 30	.219 38.653	1.00 45.36	8
ATOM	3174	OH2 WAT S 187	30.770 9	.460 69.837	1.00 32.44	8
ATOM	3175	OH2 WAT S 188	22.157 39	.535 78.736	1.00 37.01	8
MOTA	3176	OH2 WAT S 189	11.778 50	.526 68.987	1.00 41.34	8
ATOM	3177	OH2 WAT S 190	31.339 60	.910 49.439	1.00 21.88	8
ATOM	3178	OH2 WAT S 191	31.165 14	.244 74.907	1.00 27.47	8
ATOM	3179	OH2 WAT S 192	39.705 15	.398 70.464	1.00 47.05	8
ATOM	3180	OH2 WAT S 193	3.668 34	.304 72.937	1.00 39.82	8
MOTA	3181	OH2 WAT S 194	25.256 9	.360 67.925	1.00 33.21	8
ATOM	3182	OH2 WAT S 195	47.575 17	.667 48.773	1.00 40.79	8
ATOM	3183	OH2 WAT S 196	32.017 13	.045 34.633	1.00 37.00	8
MOTA	3184	OH2 WAT S 197	35.476 7	.006 64.436	1.00 49.59	' 8
MOTA	3185	OH2 WAT S 198	12.180 16	.270 56.288	1.00 47.22	8
ATOM	3186	OH2 WAT S 199	37.133 21	.226 75.963	1.00 38.59	8
ATOM	3187	OH2 WAT S 200	40.268 15	.712 48.199	1.00 39.24	8
ATOM	3188	OH2 WAT S 201	25. 1 59 17	.768 46.858	1.00 49.88	8
ATOM	3189	OH2 WAT S 202		.104 65.727	1.00 53.46	8
ATOM	3190	OH2 WAT S 203	36.741 20	.267 33.858	1.00 41.90	8
ATOM	3191	OH2 WAT S 204		.930 47.546	1.00 48.06	8
MOTA	3192	OH2 WAT S 205		.731 54.471	1.00 27.07	8
ATOM	3193	OH2 WAT S 206		.778 74.101	1.00 47.44	8
ATOM	3194	OH2 WAT S 207		.767 45.193	1.00 59.49	8
ATOM	3195	OH2 WAT S 208		.151 36.382	1.00 45.31	8
MOTA	3196	OH2 WAT S 209		.405 51.989	1.00 58.23	8
ATOM	3197	OH2 WAT S 210		.551 36.157	1.00 30.78	8
ATOM	3198	OH2 WAT S 211		.360 63.283	1.00 34.08	8
ATOM	3199	OH2 WAT S 212		.471 49.568	.1.00 33.92	8
MOTA	3200	OH2 WAT S 213		.615 33.729	1.00 44.22	8
ATOM	3201	OH2 WAT S 214		.496 75.246	1.00 40.49	8
MOTA	3202	OH2 WAT S 215		.473 46.604	1.00 39.35	8
ATOM	3203	OH2 WAT S 216		.987 44.362	1.00 36.50	8
ATOM	3204	OH2 WAT S 217		.281 65.917	1.00 34.22	8
MOTA	3205	OH2 WAT S 218	32.823 8	.977 59.213	1.00 27.03	8
END						

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						Figure 18					
					sidue		Y	Z	В	Segment	ΪD
ATOM	1		ALA		2	46.726	14.97	1 138.208	1.00	56.80	
MOTA	2		ΑLλ	A.	2	47.943	12.81	3 138.561	1.00	58.93	
ATOM	3	0	ALA	Α	2	48.857	13.292	2 137.884	1.00	60.99	
ATOM	4	N	ALA	A	2	46.995	14.04	5 140.488		56.88	
MOTA	5	CA	ALA	A	2	46.801	13.697			59.41	
ATOM-	6	N	LYS		3	47.890				53.81	
ATOM	7	CA	LYS		3	48.937	10.59			53.62	
ATOM	8		LYS		3	48.736	9.229				
MOTA	9	-	LYS		3	48.917	9.279			50.26	
	10									56.64	
ATOM			TAS		3	48.950	7.891			57.18	
ATOM	11		LYS		3	49.160		142.796		56.74	
MOTA	12	NZ	LYS		3	50.423	8.663			54.86	
MOTA	13	С	LYS		3 .	49.063		136.986	1.00	49.95	
ATOM	14	0	LYS		3	48.088	10.562		1.00	44.34	
ATOM	15	N	VAL	A	4	50.287	10.147	136.550	1.00	46.01	
ATOM	. 16	CA	VAL	A	4	50.609	9.985	135.142	1.00	42.48	
ATOM	17	CB	VAL	A	4	51.901	10.755	134.809	1.00	43.42	
ATOM	18	ÇG1	VAL	A	4	52.179	10.713		1.00	39.20	
ATOM	19	CG2	VAL	A	4	51.773		135.310		39.34	
ATOM	20	C	VAL		4	50.787	8.510	134.806		38.41	
ATOM	21	ō	VAL		4	51.659	7.839	135.351		37.08	
ATOM	22	N	LYS		5	49.959	8 011	133.899			
ATOM	23	CA	LYS		5	50.016	6.610			37.79	
ATOM	24	CB	LYS		5					38.17	
	25	CG				48.700	5.915			38.40	
MOTA	26		LYS		5	48.411	5.803			42.84	•
ATOM		CD	LYS	A	5	49.384	4.855			44.10	
ATOM	27	CE	LYS		5	49.017	4.632			45.97	
ATOM	28	NZ	LYS		5	49.045		138.322		51.78	
ATOM	29	C	LYS		5	50.275	6.392			38.31	
ATOM	30	0	LYS		5	49.992	7.253			38.13	
MOTA	31	N	LEU		6	50.817	5.220	131.717	1.00	35.05	
ATOM	32	CA	LEU .		5	51.082		130.346	1.00	31.46	
MOTA	33	CB	LEU	A	5	52.582	4.592	130.133	1.00	28.46	
MOTA	34	CG	LEU .	A	5.	53.094	4.256	128.720	1.00	30.91	
ATOM	35	CD1	LEU .	A (5	52.618	2.884	128.295	1.00	33.05	
ATOM	36	CD2	LEU .	A (5	52.630	5.312	127.744		21.96	
ATOM	37	С	LEU .	A (ŝ	50.307	3.512			30.50	
MOTA	38	0	LEU .	a (5	50.453	2.581	130.955		32.82	
ATOM	39	N	ILE .			49.459	3.456			26.94	
ATOM	40	CA	ILE :			48.676	2.255	128.893		28.29	
ATOM	41	CB	ILE			47.218	2.598			28.94	
ATOM	42	CG2	ILE			46.499	1.343			32.57	
ATOM	43	CG1	ILE			46.447	3.172	129.688			
ATOM	44	CD1	ILE A			46.979				36.59	
	45	C	ILE A				4.468			46.80	
MOTA	46					49.341	1.470	127.770	1.00		
MOTA	47	0		A 7		49.600	2.009			27.65	
MOTA						49.638	0.201		1.00		
MOTA	48	CA		A 6		50.277		127.016	1.00		
ATOM	49	C	GLY A	_		50.578		127.480	1.00		
ATOM	50	0	GLY 2			50.224		128.592	1.00		
atom	51	N	THR 3			51.238	-2.777	126.611	1.00	28.94	
MOTA	52	CA	THE ?			51.614	-4.156	126.877	1.00	33.63	
ATOM	53	CB	THR A			50.393	-5.083	126.857	1.00	36.19	
ATOM	54	OG1	THR A	A 9		50.827		126.992	1.00	34.87	
ATOM	55	CG2	THR A			49.633	-4.931	125.548	1.00	36.49	
MOTA	56	С	THR A	1 9		52.567		125.794		34.83	•
MOTA	57	0	THR A			52.545		124.677	1.00		
ATOM	58	N	LEU A			53.407		126.129	1.00		
ATOM	59	CA	LEU A			54.345		125.164	1.00	-	
	60	CB	LEU A				-0.10/	125.881			
ATOM		CG				55.402 56.482	-1.009	126.687	1.00		
ATOM	61		LEU A				~0.282	127 647	1.00		
ATOM	62		LEU A			55.870	-5.293	127.647	1.00	** .	
ATCM	63		LEU A			57.319	-7.306	127.424	1.00		
ATOM	64	C	LEU A			53.591	-7.039	124.159	1.00		
ATOM	65	0	LEU A			54.055		123.044	1.00		
MOTA	66	N	ASP A	11		52.419	-7.51 <i>9</i>	124.557	1.00	47.28	

ATOM	61	7 CA	ASP	A 11	51.617	-8.36	9 123.683	1.00	53 30
ATOM	68				50.230	-8.60	8 124.287	1.00	
ATOM	69				50.295	-9.33	1 125.610	1.00	
ATOM	70		1 ASP			-10.35	8 125.685		
MOTA	71 72				49.630		3 126.567		
ATOM ATOM	73	-	ASP ASP		51.459 51.360		0 122.257	1.00	
ATOM	74	-	TYR		51.424	-6.52	6 121.311 1 122.092		
ATOM	75				51.275		0 120.749		
ATOM	76				51.328		7 120.755	1.00	
ATOM	77	CG			50.164	-3.72	9 121.421	1.00	
MOTA	78		1 TYR		50.296	-3.15	7 122.686	1.00	
ATOM	79				49.252	-2.43		1.00	
ATOM ATOM	80 81				48.952		5 120.749	1.00 4	
ATOM	82		2 TYR . TYR .		47.906 48.061		7 121.310 9 122.566	1.00 4	
ATOM	83	OH	TYR		47.030	-1 54	3 122.366 3 123.116	1.00 4	
ATOM	84	c	TYR		52.367	-6.50	3 119.816	1.00 5	
ATOM	85	0	TYR		52.197	-6.52	118.596	1.00 4	
ATOM	86	N	GLY :	A 13	53.484		120.396	1.00 4	
MOTA	87	CA	GLY .		54.574	-7.45		1.00 5	
ATOM	. 88	C	GLY		54.196	-8.727		1.00 5	
ATOM ATOM	- 89 90	. M	GLY I		54.931 53.045	-9.184 -9.294	117.982	1.00 5	
ATOM	91	CA	LYS A				119.207 118.579	1.00 5	
ATOM	92	СВ	LYS				119.653	1.00 5	
MOTA	93	CG	LYS A		53.086	-12.062	120.591	1.00 6	2.81
ATOM	94	CD	LYS A		53.934	-13.154	119.918	1.00 6	
ATOM	95	CE	LYS A		54.747		118.734	1.00 6	
MOTA	96	NZ	LYS A			-13.713		1.00 5	
ATOM ATOM	97 98	C	LYS A			-10.231	117.559 116.942	1.00 5	
ATOM	99	Ŋ	TYR A		51.143	-R 955	117.372	1.00 5 1.00 4	
ATOM	100	CA	TYR A	-	50.091	-8:563	116.449	1.00 4	
ATOM	101	CB	TYR A	15	48.959	-7.915		1.00 5	
ATOM	102	CG	TYR A		48.456	-8.793		1.00 5	3.01
ATOM	103		TYR A		48.166	-8.255		1.00 5	
ATOM ATOM	104 105	CE1			47.722	~9.053		1.00 5	
ATOM	106	CE2				-10.166 -10.976		1.00 5 1.00 5	
ATOM	107	cz	TYR A			-10.412		1.00 5	
MOTA	108	OH	TYR A			-11.208		1.00 5	
ATOM	109	С	TYR A		50.592		115.353	1.00 4	6.20
ATOM	110	0	TYR A		49.933	-6.635		1.00 4	
MOTA	111	N CA	ARG A		51.758		114.791	1.00 4	
ATOM ATOM	113	CB	ARG A		52.3 <u>47</u> 53.779	-7.109 -7.545		1.00 49	
ATOM	114	CG	ARG A	16	54.677	-7.698		1.00 5	
ATOM	115	CD	ARG A		54.992	-6,388		1.00 60	
MOTA	116	NE	ARG A	16	56.021		116.328	1.00 6	
MOTA	117	CZ		16	57.211		116.070	1.00 6	5.68
MOTA	118		ARG A	16	57.520			1.00 69	
MOTA	119		ARG A	16	58.093		117.046	1.00 66	
MOTA MOTA	120 121	C O	ARG A	16 16	51.573 50.871		112.429 112.254	1.00 44	
ATOM	122	N	TYR A	17	51.715		111.514	1.00 43	
ATOM	123	CA	TYR A	17	51.067		110.215	1.00 38	
ATOM	124	CB	TYR A	17	50.913		109.565	1.00 33	
. MOTA	125	CG	TYR A	17	49.744		110.084	1.00 27	.35
ATOM	126	CD1	TYR A	17	49.598		111.443	1.00 27	
aton	127	CE1	TYR A	17	48.540		111.909	1 00 27	
ATOM	128 129		TYR A	17 17	48.807 47.752		109.204	1.00 25	
atom atom	130	CZ	TYR A	17	47.626		109.656 111.009	1.00 26	
atom atom	131	•	TYR A	17	46.602		111.450	1.00 27	
ATOM	132		TYR A	17	51.972		109.368	1.00 41	
-							_		

ATOM	13.	3 о	TYR	A 17		53.15	0 -7.5	25 109.68	3 1 00 25 52
MOTA	13	4 N	PRO			51.44			
MOTA	13	5 CI				50.07			
MOTA	13		PRO	A 18		52.20			,
ATOM	137					51.21			
ATOM	138					50.34			
ATOM	139	_	PRO			53.55			
MOTA	140		PRO			53.78	8 -7.10		
ATOM	141		LYS			54.432	2 -9.26		
ATOM	142					55.800	0 -9.04	4 106.114	1.00 57.00
ATOM	143					56.223			1.00 62.34
ATOM	144			A 19		55.069	9 -10.92	9 104.537	1.00 67.94
ATOM	145					54.239			1.00 70.76
MOTA	146					53.004	1 -10.65		1.00 73.70
ATOM ATOM	147		LYS			52.116		_	1.00 79.01
ATOM	148 149		LYS A			56.229			
ATOM	150		LYS A			57.230			
MOTA	151	CA	ASN A	-		55.515			1.00 49.62
ATOM	152	CB	ASN A			55.925			1.00 50.02
ATOM	153	CG	ASN A			55.829			1.00 50.62
ATOM	154	OD1				56.729			1.00 51.26
ATOM	155	ND2				57.948 56.130			1.00 46.88
ATOM	156	C	ASN A			55.167			1.00 50.85
ATOM	157	ō	ASN A			55.481	-4.86 -3.77		1.00 45.50
ATOM	158	N	HIS A			54.182	-4.99		1.00 45.35
ATOM	159	CA	HIS A			53.374	-3.86		1.00 37.46
ATOM	160	CB	HIS A			52.198	-4.35		1.00 32.39 1.00 29.34
ATOM	161	CG	HIS A	21		51.118	-3.339		1.00 30.50
ATOM	162		HIS A	21		50.999	-2.314		1.00 22.88
MOTA	163	ND1	HIS A	21		49.993	-3.298		1.00 30.15
ATOM	164		HIS A	21		49.226	-2.293		1.00 30.96
ATOM	165	NE2		21		49.814	-1.680	106.945	1.00 36.41
MOTA	166	C	HIS A	21		54.194	-2:879		1.00 29.18
ATOM	167	0	HIS A	21		55.030	-3.279		1.00 26.92
ATOM ATOM	168 169	N CD	PRO A	22		53.965	-1.572		1.00 31.12
ATOM	170	CA	PRO A	22		53.027	-0.912		1.00 29.46
ATOM	171	CB	PRO A	22 22		54.702	-0.567		1.00 29.27
ATOM	172	CG	PRO A	22		54.012 53.670	0.732		1.00 26.00
ATOM	173	·c	PRO A	22		54.624	0.434 -0.822		1.00 31.52
ATOM	174	Ō	PRO A	22		55.575	-0.538		1.00 29.96
ATOM	175	N	LEU A	23		53.501	-1.371		1.00 27.47 1.00 26.64
MOTA	176	CA	LEU A	23		53.309	-1.644		1.00 20.64
MOTA	177	CB	LEU A	23		51.833	-1.428		1.00 24.09
MOTA	178	ĆG	LEU A.	23		51.356	0.029		1.00 25.30
ATCM	179	CD1	LEU A	23 ·		49.836	0.103	110.668	1.00 17.72
MCTA	180		LEU A	23		52.086	0.816	111.574	1.00 24.15
atom	181		LEU A	23		53.775	-3.015	110.662	1.00 31.64
ATCM	182	0	LEU A	23		53.252	-3.512	111.667	1.00 31.00
ATOM	183		LYS A	24		54.753	-3.636	110.012	1.00 28.25
atom	184		LYS A	24		55.200		110.513	1.00 30.90
ATOM	185		LYS A	24		55.718	-5.810	109.372	1.00 36.59
MOTA	186	CG	LYS A	24		57.178		108.982	1.00 40.77
ATOM	187 188	CD :	LYS A	24		57.546		108.535	1.00 44.61
ATOM		CE	LYS A	24	•	58.858		107.755	1.00 50.44
ATOM	190		LYS A LYS A	24		59.959		108.487	1.00 51.30
atom Atom			LYS A	24		56.282		111.581	1.00 32.57
atom atom			ILE A	24		56.695		112.245	1.00 29.83
atum Atom			ILE A	25 25		56.729		111.750	1.00 27.06
ATOM			ILE A	25		-57.755 58.416		112.739 112.499	1.00 30.45
ATOM			ILE A	25		59.056		112.499	1.00 33.37
ATOM ·			ILE A	25		57.361		112.662	1.00 33.22
			ILE A	25		57.930		112.700	1.00 30.45 1.00 33.12
ATCM			ILE A	25		57.156		114.141	1.00 33.12
	-	_		-			3.143	T	T. OO JE . TO

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Figure 18-4

				•				
ATOM	199 O ILE	A 25		55.96	7 -2.85	1 114.310	1.00 28.1	_
MOTA	200 N PRO			57.979		2 115.168		
ATOM	201 CD PRO			59.39		8 115.139		
ATOM	202 CA PRO			57.507		2 116.556		
MOTA	203 CB PRO			58.709		0 117.347		4
ATOM	204 CG PRO			59.454		1 116.324		Ţ
MOTA	205 C PRO			57.265		0 116.827		
ATOM	206 O PRO	-		58.001		4 116.315		
ATOM	207 N ARG	-		56.251		4 117.614		٥
ATOM	208 CA ARG			55.977		5 117.899		
ATOM	209 CB ARG			54.787		3 117.039 3 117:048		
ATOM	210 CG ARG		•	55.075	0.33	117.046		7
ATOM	211 CD ARG			53.918	0.19.	3 114.620	1.00 29.6	
ATOM	212 NE ARG A			53.622		114.520	1.00 26.6	
ATOM	213 CZ ARG			52.649		115.173	1.00 28.5	2
ATOM	214 NH1 ARG A			51.857		115.173	1.00 29.7	U
ATOM	215 NH2 ARG			52.451		114.983	1.00 30.1	
MOTA	216 C ARG A			55.746	0 11/	119.387	1.00 23.2	5
ATOM	217 O ARG A			56.679		120.113	1.00 30.7	Ţ
ATOM	218 N VAL A			54.529			1.00 24.6	
ATOM	219 CA VAL A			54.282			1.00 23.5	
ATOM	220 CB VAL A			52.800			1.00 29.3	
ATOM	221 CG1 VAL A					121.635	1.00 34.50	
ATOM	222 CG2 VAL A			52.599		123.142	1.00 32.4	2
ATOM	223 C VAL A			51.947 55.158		120.903	1.00 33.7	7
ATOM	224 O VAL A			55.673		122.145	1.00 29.79	دَ
ATOM	225 N SER A			55.341		123.182 121.718	1.00 32.49	
ATOM	226 CA SER A			56.162	-2.982		1.00 26.09	
ATOM	227 CB SER A		•	56.058	-4.399		1.00 31.39	
ATOM	228 OG SER A	29		56.562		121.905° 120.579	1.00 26.92	:
ATOM	229 C SER A	29		57.609	-2.482	120.579	1.00 33.85	
ATOM	230 O SER A	29		58.378	-2.402	123.391	1.00 34.77	
ATOM	231 N LEU A	30		57.967		121.380	1.00 29.39	
ATOM	232 CA LEU A	30		59.317		121.240	1.00 31.20	
ATOM	233 CB LEU A	30		59.554		119.829	1.00 32.03	
ATOM	234 CG LEU A	30		61.008		119.333	1.00 33.22	
ATOM	235 CD1 LEU A	30		61.066		118.224	1.00 28.76	
ATOM	236 CD2 LEU A	30		61.948	-0.135	120.441	1.00 35.11	
MOTA	237 C LEU A	30		59.423		122.236	1.00 30.29	
ATOM	238 O LEU A	30		60.397	0.019	122.984	1.00 27.69	
ATOM	239 N LEU A	31		58.408		122.232	1.00 27.38	
ATOM	240 CA LEU A	31		58.372	1.915	123.126	1.00 24.94	
ATOM	241 CB LEU A	31		57.008		123.042	1.00 24.92	
ATOM	242 CG LEU A	31		56.918	4.069	123:460	1.00 30.49	
ATOM	243 CD1 LEU A	31		55.492	4.390	123.881	1.00 24.71	
MOTA	244 CD2 LEU A	31		57.851		124.603	1.00 27.32	
MOTA	245 C LEU A	31		58.610	1.429		1.00 28.18	
ATOM	246 O LEU A	31		59.489	1.928	125.263	1.00 33.64	
ATOM	247 N LEU A	32		57.831	0.445	125.000	1.00 30.17	
ATOM	248 CA LEU A	32		57.965		126.357	1.00 30.59	
MOTA	249 CB LEU A	32				126.601	1.00 30.55	
MOTA	250 CG LEU A	32		55.458		126.402	1.00 29.50	
MOTA	251 CD1 LEU A	32		54.611		126.727	1.00 28.31	
MOTA	. 252 CD2 LEU A	32		55.058	0.273	127.287	1.00 31.92	
MOTA	253 C LEU A	32		59.376		126.657	1.00 33.56	
MOTA	254 O LEU A	32		59.961		127.682	1.00 36.51	
ATOM	255 N ARG A	33		59.926		125.777	1.00.29.75	
ATOM	256 CA ARG A	33 -		61.271		125.999	1.00 33.49	
ATOM	257 CB ARG A	33		61.630		124.945	1.00 39.50	
ATOM	258 CG ARG A	33		60.814		125.024	1.00 44.40	
ATOM	259 CD ARG A	33		61.237	-5.256		1.00 53.68	
ATOM	260 NE ARG A	33		60.515	-6.522		1.00 56.66	
ATOM	261 CZ ARG A	33		60.611	-7.384		1.00 58.73	
ATOM	262 NH1 ARG A	33		61.402	-7.121	126.045	1.00 59.32	
ATOM	263 NH2 ARG A	33		59.911	-8.511	124.991	1.00 57.91	
ATOM	264 C ARG A	33	•	62.314	-0.845		1.00 31.45	
					0.010			

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Figure 18-5

MOTA	265 O ARG A 33		63.288	0.005 105 ===
ATOM				
ATOM	200 122 1		62.103	0.146 125.123 1.00 32.4
	and a ja		63.042	1.253 125.000 1.00 33.3
ATOM	268 CB PHE A 34		62.617	1.00 33.3
ATOM	269 CG PHE A 34			
ATOM	270 CD1 PHE = 34		63.653	
	- · · · · · · · · · · · · · · · · · · ·		64.825	2.819 122.838 1.00 29.21
ATOM	271 CD2 PHE A 34.		63.458	
MOTA	272 CE1 PHE A 34		65.793	
ATOM	273 CE2 PHE A. 34			
ATOM	274 CZ PHE A 34		64.416	5.501 123.435 1.00 32.67
	: n 34		65.589	5.108 122.783 1.00 29.08
ATOM	275 C PHE A 34		63.083	
ATOM	276 O PHE A 34		64.155	
ATOM	000			2.294 126.852 1.00 27.49
ATOM	27/ N LYS A 35 278 CA LYS A 35		61.912	2.432 126.802 1.00 29.34
			61.848	3.191 128.042 1.00 31.48
ATOM	279 CB LYS A 35		60.406	
ATOM	'280 CG LYS A 35			
ATOM			59.803	4.552 127.395 1.00 32.98
			58.404	4.974 127.790 1.00 40.93
ATOM	282 CE LYS A 35		57.410	3.827 127.688 1.00 44.56
ATOM	283 NZ LYS A 35		57.754	00 44.50
ATOM	284 C LYS A 35			2.50 33.10
ATOM	55		62.443	2.387 129.183 1.00 34.47
	204		63.136	2.933 130.043 1.00 32.01
MOTA	286 N ASP A 36		62.180	
ATOM	287 CA ASP A 36		62.710	
ATOM	288 CB ASP A 36		62.710	2.00 37.33
ATOM			62.145	-1.178 130.126 1.00 41.27
			62.731	-2.117 131.157 1.00 43 77
ATOM	290 OD1 ASP A 36		62.660	-1.793 132.360 1.00 43.92
ATOM	291 OD2 ASP A 36		63.261	
ATOM	292 C ASP A 36		64.227	
ATOM	293 O ASP A 36			0.181 130.174 1.00 38.74
ATOM			64.902	0.187 131.201 1.00 36.23
	294 N ALA A 37		64.760	0.127 128.958 1.00 37.96
MOTA	295 CA ALA A 37	•	66.201	
ATOM	296 CB ALA A 37		66.525	
MOTA	297 C ALA A 37			
ATOM			66.832	1.386 129.244 1.00 40.09
			67.962	1:402 129.714 1.00 38.80
MOTA	299 N MET A 38		66.085	2.477 129.131 1.00 39.04
ATOM	300 CA MET A 38		66.567	
ATOM	301 CB MET A 38		65.965	4 060 400 410
ATOM	200		05.905	4.863 128.640 1.00 36.66
	2.2.2		66.335	4.744 127.173 1.00 39.16
ATOM	303 SD MET A 38		68.005	5.298 126.840 1.00 37.55
ATOM	304 CE MET A 38		67.892	7 000 100 000
ATOM	305 C MET A 38		66.187	
ATOM	306 O MET A 38			4.094 130.995 1.00 40.58
ATOM			66.484	5.173 131.502 1.00 38.12
			65.530	3.147 131.657 1.00 38.41
MOTA	308 CA ASN A 39		65.094	3.346 133.039 1.00 42.46
ATOM	309 CB ASN A 39		66.298	
ATOM	310 CG ASN A 39		67.125	
ATOM	311 OD1 ASN A 39			2 224 134.074 1.00 51.69
ATOM			66.625	2.175 134.487 1.00 54.33
	312 ND2 ASN A 39		68.396	2 313 133.695 1.00 49.13
ATOM	313 C ASN A 39		64.222	4.594 133.134 1.00 41.19
ATOM	314 O ASN A 39		64.375	
ATOM	315 N LEU A 40			5.402 134.050 1.00 42.74
ATOM			63.301	4.746 132.188 1.00 40.22
			62.427	5.909 132.170 1.00 39.85
ATOM	317 CB LEU A 40		62.524	6.610 130.812 1.00 40.42
ATOM	318 CG LEU A 40		63.940	
ATOM ·	319 CD1 LEU A 40			
ATOM			63.916	7.753 129.088 1.00 32.75
			64.470	8.031 131.513 1.00 38.89
ATOM	321 C LEU A 40		60.967	5.610 132.505 1.00 38.97
ATOM	322 O LEU A 40		60.076	
ATOM	323 N ILE A 41			
ATOM			50.720	4.461 133.124 1.00 38.57
		;	59.363	4.109 133.520 1.00 42.43
atom	325 CB ILE A 41	5	58.536	3.575 132.330 1.00 39.13
ATOM	326 CG2 ILE A 41			
ATOM	327 CG1 ILE A 41	2		
ATOM				3.367 132.774 1.00 38.71
			6.147	2.920 131.676 1.00 44.09
MOTA	329 C ILE A 41		9.376	3.056 134.619 1.00 42.40
ATCM .	330 O ILE A 41			
•				2.195 134.654 1.00 43.05

ATOM	331 N ASP A 42	58.414 3.148 135.532 1 00 47 83
ATOM		50 301 2 107 106 11.00 47.00
ATOM		58.301 2.183 136.620 1.00 49.23 58.243 2.880 137 984 1 00 46 66
MOTA		50 402 2 600 400 201
MOTA		60 614
ATOM	336 OD2 ASP A 42	60.614 3.146 138.141 1.00 49.28
ATOM		59.355 4.866 138.678 1.00 52.47
ATOM		57.034 1.368 136.405 1.00 51.22
ATOM	220 11	56.048 1.866 135.864 1.00 48.07
MOTA	340	57.072 0.111 136.832 1.00 51.41
ATOM	244	55.945 -0.792 136.673 1.00 50.67
ATOM	240	56.234 -2.094 137.412 1.00 54.49
ATOM	242	55.208 -3.178 137.185 1.00 60.55
ATOM	344 ODO N 43	55.524 -4.432 137.974 1.00 66.12
ATOM	. 246	54.761 -5.417 137.861 1.00 70.33
MOTA	246	56.536 -4.427 138.711 1.00 67.39
ATOM	7.5	54.645 -0.178 137.178 1.00 50.20
ATOM	210	53.567 -0.475 136.658 1.00 48.27
ATOM		54.755 0.683 138.186 1.00 49.04
ATOM	250	33.601 1.356 138.778 1.00 47.56
ATOM	200	54.013 2.004 140.112 1.00 54.73
ATOM	750	53.190 3.229 140.542 1.00 58.07
ATOM		53.705 4.500 139.853 1.00 61 24
ATOM		52.849 5.727 140.151 1.00 61.93
ATOM	250 -	51.501 5.644 139.519 1.00 62.80
ATOM	356 2	52.929 2.387 137.875 1.00 44.52
ATOM		51.752 2.701 138.052 1.00 45 31.
ATOM	350	53.674 2.915 136.914 1.00 41.03
ATOM	250	53.140 3.914 135.994 1.00 41.23
MOTA	260	54.271 4.810 135.500 1.00 38.52
ATOM	7.64	54.973 5.572 136.589 1.00 40.30
ATOM	250	56.241 6.222 136.096 1.00 38.06
ATOM	242	57.170 5.478 135.715 1.00 36.93
ATOM		56.306 7.467 136.084 1.00 32.57
ATOM	364 C GLUA 45. . 365 O GLUA 45	52.479 3:253 134.791 1.00 40.69
ATOM	366 N LEUA 46	51.783 3.907 134.015 1.00 39.77
ATOM	367 CA LEU A 46	52.700 1.953 134.645 1.00 36.90 52.165 1.207 133.517 1.00 40 46
ATOM	368 CB LEU A 46	E3 333 0 010 111
ATOM ·	369 CG LEU A 46	50 077
ATOM	370 CD1 LEU A 46	53 F71 0 200 100 43.73
ATOM	371 CD2 LEU A 46	5.4 035
ATOM	372 C LEU A 46	50 953
ATOM	373 O LEU A 46	50 741
ATOM	374 N ILE A 47	10 051
ATOM	375 CA ILE A 47	49 500 0 000 100 000 100
ATOM	376 CB ILE A 47	10 32.12
ATOM	377 CG2 ILE A 47	16 060
Arom	378 CG1 ILE A 47	1 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2
ATOM	379 CD1 ILE A 47	1.00 50.50
ATOM	380 C ILE A 47	10 100 35.40
ATOM	381 O ILE A 47	100 33.07
ATOM	382 N LYS A 48	40.000
ATOM	383 CA LYS A: 48	
MOTA	384 CB LYS A 48	10 100
MOTA	385 CG LYS A - 48	
MOTA	386 CD LYS A 48	
MOTA	387 CE LYS A 48	47 067
MOTA	388 NZ LYS A 48	
MOTA	389 C LYS A 48	
TCM	390 C LYS A 48	15 364
TOM	391 N SER A 49	17 071
TOM	392 CA SER A 49	45 000
MOT	393 CB SER A 49	
TOM	394 OG SER A 49	47 571
TOM	395 C SER A 49	1.00 30.74
TOM	396 O SER A 49	
		45.295 -5.059 128.436 1.00 34.44

ATOM	397 N ARG A 50	43.688 -3.582 127.922 1.00 32.87
ATOM	398 CA ABG A 50	42.632 -4.582 127.960 1.00 31.45
MOTA	399 CB ARG A 50	41.636 -4.325 129.101 1.00 28.35
ATOM	400 CG ARG A 50	40.729 -3.103 128.915 1.00 32.05
ATOM	401 CD ARG A 50	39.653 -3.055 130.008 1.00 30.46
ATOM	402 NE ARG A 50	38.821 -1.850 129.964 1.00 25.21
ATOM	403 CZ ARG A 50	37.930 -1.569 129.016 1.00 28.32
ATOM	404 NH1 ARG A 50	37.726 -2.406 128.001 1.00 25 45
MOTA	405 NH2 ARG A 50	37.238 -0.439 129.087 1.00 24.92
MOTA	406 C ARG A 50	41.894 -4.470 126.638 1.00 31.12
ATOM	407 O ARG A 50	41.895 -3.406 126.019 1.00 24.62
ATOM	408 N PRO A 51	41.264 -5.566 126.181 1.00 32.55
MOTA	409 CD PRO A 51	41.164 -6.921 126.751 1.00 32.40
MOTA	410 CA PRO A 51	40.534 -5.506 124.917 1.00 30 36
ATOM	411 CB PRO A 51	40.138 -6.967 124.683 1.00 33.95
ATOM	412 CG PRO A 51	41.173 -7.750 125.499 1.00 32.85
ATOM	413 C PRO A 51	39.309 -4.630 125.134 1.00 31.61
ATOM	414 O PRO A 51	38.877 -4.431 126.267 1.00 29.84
ATOM	415 N ALA A 52	38.755 -4.093 124.058 1.00 29.09
MOTA	416 CA ALA A 52	37.556 -3.294 124.183 1.00 29.61
ATOM	417 CB ALA A 52	37.365 -2.447 122.956 1.00 28.67
ATOM	418 C ALA A 52	36.437 -4.321 124.288 1.00 32.39
ATOM	419 O ALA A 52	36.603 -5.453 123.844 1.00 30.40
ATOM	420 N THR A 53	35.318 -3.947 124.896 1.00 32.98
ATOM ATOM	421 CA THR A 53 422 CB THR A 53	34.192 -4.868 124.997 1.00 36.61
ATOM	422 CB THR A 53 423 OG1 THR A 53	33.253 -4.514 126.166 1.00 34.22
ATOM	424 CG2 THR A 53	32.734 -3.193 125.970 1.00 29.52 33.998 -4.579 127.493 1.00 36.45
ATOM	425 C THR A 53	
ATOM	426 O THR A 53	
ATOM	427 N LYS A 54	1.00 32.07
ATOM	428 CA LYS A 54	
MOTA	429 CB LYS A 54	
ATOM	430 CG LYS A 54	30.933 -6.851 121.994 1.00 41.68 30.367 -7.034 120.597 1.00 49.42
ATOM	431 CD LYS A 54	29.541 -8.310 120.508 1.00 51.82
ATOM	432 CE LYS A 54	29.075 -8.588 119.087 1.00 52.94
ATOM	433 NZ LYS A 54	30.216 -8.879 118.182 1.00 54.26
ATOM	434 C LYS A 54	30.913 -4.347 122.237 1.00 39.46
MOTA	435 O LYS A 54	30.719 -3.637 121.249 1.00 37.19
ATOM	436 N GLU A 55	30.404 -4.075 123.434 1.00 36.71
ATOM	437 CA GLU A 55	29.554 -2.913 123.665 1.00 36.18
ATOM	438 CB GLU A 55	29.109 -2.877 125.127 1.00 42.16
MOTA	439 CG GLU A 55	28.223 -1.694 125.476 1.00 46.04
ATOM ATOM	440 CD GLUA 55 441 OE1 GLUA 55	27.873 -1.639 126.953 1.00 51.15
	441 OE1 GLU A 55 442 OE2 GLU A 55	27.092 -0.748 127.343 1.00 56.53
ATOM ATOM	443 C GLUA 55	28.382 -2.482 127.727 1.00 51.67
ATOM	444 O GLUA 55	30.278 -1.607 123.32: 1.00 35.45 29.721 -0.729 122.66 1.00 29.11
ATOM	445 N GLUA 56	
ATOM	446 CA GLU A 56	20 000
ATOM	447 CB GLU A 56	22 (27 0 222 224 224
ATOM	448 CG GLU A 56	33.635 -0.329 124.232 1.00 30.71 33.474 -0.484 125.746 1.00 35.09
MOTA	449 CD GLU A 56	34.787 -0.675 126.479 1.00 32.29
ATOM	450 OE1 GLU A 56	35.645 -1.434 125.986 1.00 34.54
MOTA	451 OE2 GLU A 56	34.951 -0.094 127.569 1.00 33.25
ATOM	452 C GLU A 56	32.495 -0.104 121.988 1.00 32.51
ATOM	453 O GLUA 56	32.341 0.990 121.444 1.00 29.59
ATOM	454 N LEUA 57	32.827 -1.196 121.311 1.00 35.58
ATOM	455 CA LEU A 57	33.039 -1.147 119.871 1.00 35.70
MOTA	456 CB LEU A 57	33.475 -2.512 119.334 1.00 35.25
ATOM	457 CG LEU A 57	34.829 -3.030 119.814 1.00 36.19
ATOM	458 CD1 LEU A 57	35.095 -4.390 119.183 1.00 33.69
MOTA	459 CD2 LEU A 57	35.925 -2.041 119.433 1.00 32.61
ATOM	460 C LEU A 57	31.772 -0.717 119.157 1.00 36.02
ATOM	461 0 LEU A 57	31.828 0.067 118.205 1.00 32.72
ATCM	462 N LEU A 58	30.631 -1.228 119.620 1.00 32.35

ATON	463 CA LEU A 58	20 750
ATON	DEG 12 JO	29.353 -0.898 119.004 1.00 33.21
ATOM	1 165 22 22 23	28.260 -1.844 119.495 1.00 35.17
ATOM	466 and 100 A 10	28.504 -3.296 119.077 1.00 33 71
ATOM	467 000 000	27.338 -4.166 119.524 1.00 36 80
ATOM	150	28.665 -3.364 117.570 1:00 36.50
	1 150 0	28.940 0.543 119.222 1.00 30.99
MOTA		27.915 0.985 118.700 1.00 35.50
ATOM	- 220 A 33	29.733 . 1.279 119.993 1 00 32 55
ATOM		29.443 2.687 120.217 1 00 30 37
ATOM		30.387 3.279 121.268 1.00 28.01
ATOM	HAU A	30.174 2.828 122.716 1.00 32.19
ATOM		31.248 3.427 123.604 1.00 24.85
ATOM		28.785 3.263 123.192 1.00 25.65
ATOM	au	29.632 3.405 118.890 1.00 31.26
ATOM		29.020 4.442 118.652 1.00 31.80
MOTA	478 N. PHE A 60	1.00 31.00
ATOM	479 CA PHE A 60	
MOTA	480 CB PHE A 60	1.00 30.24
MOTA	481 CG PHE A 60	
ATOM	482 CD1 PHE A 60	
ATOM	483 CD2 PHE A 60	
ATOM	484 CE1 PHE A 60	
MOTA	485 CE2 PHE A 60	1.00 20.12
ATOM	486 CZ PHE A 60	120 120 1.00 23.98
ATOM	487 C PHE A 60	
ATOM	488 O PHE A 60	20 010 2 054 114 555
ATOM	489 N HIS A 61	
MOTA	490 CA HIS A 61	71 075 21.00 32.65
ATOM	491 CB HIS A 61	31.075
ATOM	492 CG HIS A 61	33.576 0.238 114.116 1.00 34.25
ATOM	493 CD2 HIS A 61	34.225 0.532 112.967 1.00 34.67
ATOM ATOM	494 ND1 HIS A 61	34.328 0.786 115.133 1.00 37.78
ATOM	495 CE1 HIS A 61 496 NE2 HIS A 61	35.390 1.382 114.619 1.00 37.50
ATOM	400	35.350 1:243 113.307 1.00 37 91
ATOM	444	29.824 -0.449 114.480 1.00 38 44
ATOM	498 0 HIS A 61 499 N THR A 62	29.213 -0.612 115.538 1.00 35.78
ATOM	500 CA THR A 62	29.462 -1.015 113.327 1.00 39.73
ATOM	501 CB THR A 62	28.278 -1.868 113.218 1.00 38.05 27.682 -1.825 111.804 1.00 37.22
ATOM	502 OG1 THR A 62	29 631 2 745 110 37.22
MOTA	503 CG2 THR A 62	27 240 0 404
ATOM	504 C THR A 62	20 500
ATOM	· 505 0 THR A 62	70 777 7 750 440
ATOM	506 N GLUA 63	27 502 4 024 414 22
ATOM	507 CA GLU A 63	27 606 5 444 474 474
MOTA	508 CB GLU A 63	26.00
ATOM	509 CG GLU A 63	26.303 -6.000 114.704 1.00 43.19 26.269 -7.451 115.171 1.00 46.90
ATOM	510 CD GLU A 63	26.472 -7.593 116.665 1.00 53.11
MOTA	511 OE1 GLU A 63	26.601 -8.739 117.152 1.00 52.78
ATOM	512 OE2 GLU A 63	26.487 -6.556 117.358 1.00 57.24
ATOM	513 C GLU A 63	28.320 -6.263 113.268 1.00 36 19
ATOM	514 O GLU A 63	29.272 -7.011 113.481 1.00 29.70
MOTA	515 N ASP A 64	27.755 -6.119 112.074 1.00 35.85
ATOM	516 CA ASP A 64	28.198 -6.841 110.886 1.00 37.61
ATOM	517 CB ASP A 64	27.363 -6.382 109.697 1.00 43.30
MOTA	518 CG ASP A 64	27.313 -4.872 109.582 1.00 53.38
MOTA MOTA	519 OD1 ASP A 64 520 OD2 ASP A 64	28.290 -4.269 109.089 1.00 52.15
ATOM		26.298 -4.285 110.018 1.00 53.97
ATOM	521 C ASP A 64 522 D ASP A 64	29.673 -6.660 110,594 1.00 35.04
ATOM	523 N TYR A 65	30.379 -7.625 110.303 1.00 33.60
ATOM	524 CA TYR A 65	30.144 -5.423 110.671 1.00 33.88
ATCM	525 CB TYR A 65	31.554 -5.153 110.419 1.00 32.91 31.793 -3.637 110 375 1 00 34 80
ATOM	526 CG TYR A 65	22 242
ATOM	527 CD1 TYR A 65	34 000 33.13
ATOM	528 CE1 TYR A 65	35 35
		35.352 -3.411 109.024 1.00 32.52

ATOM	529	CD2 TYR	4 65	33.863 -2.398 111.134 1.00 34.08
ATOM	530	CE2 TYR	A 65	35.211 -2.050 111.002 1.00 29.89
MOTA	531			35.949 -2.560 109.948 1.00 35.29
MOTA	532			22 224
ATOM	533			37.286 -2.231 109.825 1.00 29.81 32.405 -5.813 111.504 1.00 27.65
ATOM	534			
ATOM	535			33.339 -6.557 111.209 1.00 27.65
				32.070 -5.559 112.765 1.00 27.32
ATOM	536			32.822 -6.153 113.858 1.00 25.82
MOTA	537			32.227 -5.764 115.217 1.00 32.25
MOTA	538		66	33.029 -6.403 116.338 1.00 28.85
ATOM	539			32.226 -4.242 115.364 1.00 31.48
ATOM	540	CD1 ILE A	66	33.607 -3.612 115.282 1.00 38.02
MOTA	541	C ILE A	66	32.836 -7.677 113.736 1.00 31.21
ATOM	542	O ILE A	66	33.891 -8.305 113.844 1.00 30.25
ATOM	543			31.672 -8.279 113.507 1.00 33.28
ATOM	544	CA ASN A		
ATOM	545		-	
ATOM	546	CG ASN A		
ATOM	547			29.338 -10.072 114.421 1.00 37.34
				29.807 -10:296 115.535 1.00 35.20
ATOM	548	ND2 ASN A		28.071 -9.709 114.236 1.00 34.83
ATOM	549	C ASN A		32.499 -10.198 112.219 1.00 31.00
ATOM	550	O ASN A		33.132 -11.248 112.306 1.00 37.26
ATOM	551	N THR A	68	32.543 -9.426 111.140 1.00 30.91
MOTA	552	CA THR A	68	33.368 -9.814 109.997 1.00 31.04
ATOM	553	CB THR A	68	33.133 -8.894 108.792 1.00 34.01
ATOM	554	OG1 THR A	68	31.780 -9.037 108.352 1.00 33.26
ATOM	.555	CG2 THR A	68	34.072 -9.256 107.646 1.00 30.84
ATOM	556	C THR A	68	· 34.844 -9.794 110.378 1.00 33.31
MOTA	557	O THR A	68	35.591 -10.708 110.024 1.00 32.52
ATOM	558	N LEU A	69	35.267 -8.768 111.117 1.00 30.30
ATOM	559	CA LEU A	69	
ATOM	560	CB LEU A	69	
ATOM	561	CG LEU A	69	
ATOM	562	CD1 LEU A	69	
ATOM	563	CD2 LEU A		37.154 -4.929 112.647 1.00 31.08
			69	37.868 -6.004 110.505 1.00 27.85
ATOM	564	C LEU A	69	37.036 -9.902 112.372 1.00 31.65
ATOM	565	O LEU A	69	38.084 -10.519 112.165 1.00 23.95
ATOM	566	N MET A	70	36.169 -10.243 113.321 1.00 30.78
ATOM	567	CA MET A	70	36.411 -11.383 114.193 1.00 34.50
MOTA	. 568	CB MET A	70	35.318 -11.486 115.258 1.00 31.96
ATOM	569	CG MET A	70	35.203 -10.259 116.147 1.00 36.26
ATOM'	570	SD MET A	70	33.948 -10.454 117.431 1.00 37.52
ATOM	571	CE MET A	70	34.633 -11.815 118.403 1.00 37.36
ATOM	572	C MET A	70	36.484 -12.685 113.401 1.00 33.33
ATOM	573	O MET A	70	37.392 -13.488 113.607 1.00 31.47
ATOM	574	N GLU A	71	35.534 -12.887 112.494 1.00 35.37
ATOM	575	CA GLU A	71	35.516 -14.098 111.681 1.00 36.6
ATOM	576	CB GLU A	71	34.245 -14.160 110.834 1.00 37.3
ATOM	577	CG GLU A	71	34.206 -15.359 109.897 1.00 46.37
ATOM	578	CD GLU A	71	
ATOM	579	OE1 GLU A	71	
		OE2 GLU A	-	.34.355 -17.733 109.952 1.00 48.94
ATOM	580		71	34.190 -16.705 111.882 1.00 45.53
MOTA	581	C GLU A	71	36.732 -14.169 110.769 1.00 35.96
ATOM	582	O GLU A	71	37.342 -15.228 110.617 1.00 32.99
ATOM	583	N ALA A	72	37.079 -13.039 110.159 1.00 36.50
MOTA	584	CA ALA A	72	38.225 -12.981 109.264 1.00 33.98
ATOM	585	CB ALA A	72	38.366 -11.580 108.675 1.00 33.23
ATOM	586	C ALA A	72	39.498 -13.362 109.998 1.00 34.60
ATOM	587	O ALA A	72	40.337 -14.094 109.466 1.00 31.53
ATOM	588	N GLU A	73	39.647 -12.873 111.224 1.00 30.87
ATOM	589	CA GLU A	73	40.847 -13.177 111.985 1.00 29.66
ATOM	590	CB GLU A	73	41.004 -12.224 113.180 1.00 33.33
ATOM	591	CG GLU A	73	42.234 -12.545 114.033 1.00 32.80
ATOM	592	CD GLU A	73	42.390 -11.634 115.233 1.00 40.07
	593	OE1 GLU A	73	42 601 - 10 410 11E 044 - 100 41 00
ATOM				42.601 -10.418 115.044 1.00 41.02
MOTA	594	OE2 GLU A	73	42.298 -12.138 116.372 1.00 41.21

MOTA	595	С	GLU A	73	40.906 -14.615 112.485 1.00 31.73
ATOM	596	0	GLU A	73	41.957 -15.249 112.409 1.00 32.96
MOTA	597	N	ARG A	74	39.798 -15.145 112.992 1.00 35.85
MOTA	598	CA	ARG A	74	
MOTA	599	CB	ARG A	74	38.548 -16.892 114.216 1.00 43.63
ATOM	600	CG	ARG A	74	37.450 -17.349 113.294 1.00 51.20
MOTA	601	CD	ARG A	74	36.366 -18.087 114.063 1.00 51.13
ATOM	602	NE	ARG A	74	35.534 -18.871 113.158 1.00 57.40
MOTA	603	CZ	ARG A	. 74	35.991 -19.870 112.403 1.00 56.36
	604		ARG A	74	37.273 -20.208 112.446 1.00 51.10
ATOM					
ATOM	605		2 ARG A	74	
MOTA	606	С	ARG A	74	40.125 -17.506 112.372 1.00 43.06
MOTA	607	0	ARG A	74	40.916 -18.429 112.541 1.00 42.52
MOTA	608	N	SER A	75	39.485 -17.305 111.222 1.00 43.63
ATOM	609	CA	SER A	75	39.670 -18.186 110.066 1.00 44.93
MOTA	610	CB	SER A	75	38.485 -18.089 109.113 1.00 42.05
	611	OG	SER A	75	38.420 -16.799 108.532 1.00 38.43
ATOM					
MOTA	612	С	SER A	75	40.910 -17.797 109.282 1.00 46.44
MOTA	613	0	SER A	75	41.339 -18.522 108.383 1.00 45.17
ATOM	614	N	GLN A	76	41.466 -16.638 109.618 1.00 46.18
MOTA	615	CA	GLN A	76	42.642 -16.116 108.936 1.00 44.73
ATOM	616	CB	GLN A	76	43.868 -16.973 109.226 1.00 37.36
ATOM	617	CG	GLN A	76	45.162 -16.208 109.045 1.00 43.96
ATOM	618	CD	GLN A	76	45.415 -15.214 110.176 1.00 41.86
	619		L GLN A	76	44.499 -14.537 110.655 1.00 37.78
ATOM					46.669 -15.111 110.591 1.00 45.00
MOTA	620	NE2		76	
MOTK	621	С	GLN A	76	42.374 -16.120 107.429 1.00 44.17
MOTA	622	0	GLN A	76	43.233 -16.495 106.630 1.00 40.49
ATOM	623	N	SER A	. 77	41.168 -15.713 107.053 1.00 43.11
MOTA	624	CA	SER A	77	40.784 -15.667 105.649 1.00 44.66
ATOM	625	CB	SER A	77	40.182 -17.004 105.220 1.00 44.56
ATOM	626	OG	SER A	77	38.974 -17.246 105.925 1.00 42.58
ATOM	627	c	SER A	77	39.747 -14.573 105.448 1.00 44.80
				77	39.096 -14.142 106.395 1.00 45.11
MOTA	628	0	SER A		
MOTA	629	N	VAL A	78	39.590 -14.137 104.207 1.00 46.06
MOTA	630	CA	VAL A	78	38.632 -13.095 103.888 1.00 47.65
ATOM	631	CB	VAL A	78	39.107 -12.245 102.701 1.00 49.63
MOTA	632	CG1	. VAL A	78	38.076 -11.167 102.391 1.00 51.25
ATOM	633	CG2	VAL A	78	40.454 -11.627 103.017 1.00 53.00
MOTA	634	С	VAL A	78	37.275 -13.682 103.530 1.00 48.07
MOTA	635	ō	VAL A	78	37.111 -14.301 102.480 1.00 42.31
	636	N	PRO A	79	36.282 -13.492 104.407 1.00 49.82
ATOM				79	36.347 -12.782 105.696 1.00 50.81
ATOM	637	CD	PRO A		
MOTA	638	CA	PRO A	79	34.927 -13.998 104.186 1.00 51.31
ATOM	639	CB	PRO A	79	34.170 -13.450 105.396 1.00 53.13
MOTA	640	CG	PRO A	-9	35.244 -13.469 106.469 1.00 53.50
MOTA	641	С	PRO A	∵9	34.343 -13.517 102.858 1.00 52.42
MOTA	642	0	PRO A	∵9	34.670 -12.428 102.382 1.00 55.73
ATOM	643	N	LYS A	80	33.482 -14.343 102.273 1.00 49.63
ATOM	644	CA	LYS A	80	32.824 -14.053 101.002 1.00 51.62
	645	CB	LYS A	80	31.632 -15.004 100.822 1.00 53.92
MOTA					30.817 -14.808 99.545 1.00 56.27
ATOM	646	CG	LYS A	80	
MOTA	647	CD	LYS A	80	
MOTA	648	CE	LYS A	80	28.744 -15.579 98.298 1.00 56.04
ATOM	649	NZ	LYS A	80	29.471 -16.036 97.081 1.00 58.90
ATOM	650	С	LYS A	80	32.338 -12.607 100.874 1.00 51.10
ATOM	651	ō	LYS A	80	31.539 -12.140 101.689 1.00-49.22
	652	N	GLY A	81	32.821 -11.914 99.842 1.00 51.14
ATOM				81	32.418 -10.537 99.592 1.00 47.07
ATOM	653	CA	GLY A		
ATOM	654	C	GLY A	81	
ATOM	655	0	GLY A	81	32.671 -8.301 100.397 1.00 43.90
ATOM	656	N	ALA A	82	33.504 -9.942 101.681 1.00 44.50
ATOM	657	CA	ALA A	82	33.973 -9.029 102.715 1.00 44.69
ATOM	658	СВ	ALA A	82	34.497 -9.825 103.903 1.00 44.62
	659	c	ALA A	82	. 35.049 -8.073 102.215 1.00 41.82
MOTE		0	ALA A	82	35.132 -6.925 102.662 1.00 35.92
ATOM	660	J	why v	~~	

MOTA	661 N ARG A 83	35.874 -8.549 101 289 1 00 42 26
ATOM	662 CA ARG A 83	35 050 7 702.203 1.00 43.30
ATOM	553	36.959 -7.742 100.741 1.00 43.25
ATOM	05	37.715 -8.533 99.677 1.00 46.60
	CCT 1210 11 03	38.988 -7.865 99.222 1.00 51 32
ATOM	665 CD ARG A 83	30 636 0 630 00
MOTA	666 NE ARGA 83	40 000 -
ATOM	667 CZ ARG A 83	41 330 C 000 00 00 1.00 04.08
ATOM	668 NH1 ARG A 83	
ATOM	550	40.403 -5.954 97.504 1.00 62.76
	CD0 -	42.599 -6.600 97.304 1.00 59 66
ATOM	670 C ARG A 83	36.453 -6.435 100.134 1.00 44.58
ATOM	671 O ARGA 83	
ATOM	672 N GLUA 84	75 404
MOTA	673 CA GLU A 84	74 004
MOTA	674 CB GLU A 84	24 44
ATOM	CDC at	34.145 -5.765 97.367 1.00 46.27
	CD C	33.621 -7.185 97.388 1.00 52 61
MOTA	676 CD GLUA 84	34.749 -8.198 97.308 1.00 54.12
MOTA	677 OE1 GLU A 84	74 2.00 54.12
ATOM	678 OE2 GLU A 84	25 200 27.704 1.00 39.06
MOTA	679 C GLU A 84	77 774
ATOM	680 O GLU A 84	22 22 27.36
MOTA		33.692 -3.379 99.416 1.00 34.30
		33.138 -5.301 100.427 1.00 36 00
MOTA	682 CA LYS A 85	32.154 -4.646 101.280 1.00 36.95
MOTA	683 CB LYS A 85	
ATOM	684 CG LYS A 85	20 100 30.00
ATOM	685 CD LYS A 85	2.00 40.72
ATOM	686 CE LYS A 85	27 070
ATOM	CO.	25 22 2
ATOM	600	26.859 -6.513 104.287 1.00 52.72
ATOM	00	32.785 -4.008 102.513 1.00 36 48
	689 O LYS A 85	32.353 -2.949 102.966 1.00 32 97
ATOM	690 N TYR A 86	33.819 -4.649 103.041 1.00 33.69
ATOM	691 CA TYR A 86	124 460
ATOM	692 CB TYR A 86	74 410 5 201 105 201
ATOM	693 CG TYR A 86	
ATOM		32.990 -5.665 105.680 1.00 35.09
ATOM	COC	32.165 -4.765 106.351 1.00 34.06
		30.866 -5.100 106.704 1.00 34.32
ATOM	696 CD2 TYR A 86	32.470 -6.923 105.365 1.00 33.17
MOTA	697 CE2 TYR A 86	31.162 -7.271 105.716 1.00 33.91
MOTA	698 CZ TYR A 86	30 360 6 350 400
ATOM	699 OH TYR A 86	20 070 6 650 454
MOTA	700 C TYR A 86	35 001
ATOM		
ATOM	200	36.552 -3.208 104.984 1.00 36.06
	700	36.382 -3.777 102.814 1.00 36.46
MOTA	703 CA ASN A 87	37.712 -3.313 102.441 1.00 32 71
MOTA	704 CB ASN A 87	37.768 -1.791 102.576 1.00 36.26
ATOM	705 CG ASN A 87	
ATOM	706 OD1 ASN A 87	
ATOM	707 ND2 ASN A 87	
ATOM	700	2.00 45.25
ATOM		38.855 -3.956 103.217 1.00 34.73
	710	39.868 -3.315 103.512 1.00 33.23
ATOM	710 N ILE A 88	38.687 -5.237 103.523 1.00 32.48
ATOM	711 CA ILE A 88	70 676 6 000 000 000
ATOM	712 CB ILE A 88	
ATOM	713 CG2 ILE A 88	
ATOM		
		38.536 -5.707 106.461 1.00 40.45
ATOM		39.641 ~4.953 107.124 1.00 42.25
ATOM	716 C ILE A 88	40.251 -7.090 103.318 1.00 37.36
MOTA	717 O ILE A 88	39.555 -7.587 102.431 1.00 35.47
MOTA	718 N GLY A 89	1.00 33.47
ATOM	719 CA GLY A: 89	
ATOM	720 C GLY A 89	
		43.134 -7.994 101.675 1.00 34.50
MOTA		43.951 -8.777 101.186 1.00 32.81
ATOM	722 N GLY A 90	43.071 -6.710 101.335 1.00 31.39
atom	723 CA GLY A 90	44.005 -6.158 100.371 1.00 23.90
ATOM	724 C GLY A 90	45.340 -5.893 101.040 1.00 28.78
MOTA	725 O GLY A 90	1.00 28.78
ATOM	726 N TYR A 91	
		46.221 -5.155 100.367 1.00 28.26

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Figure 18-12

MOTA	727 CA TYR A 91	47.539 -4.850 100 918 1 00 27 34
MOTA	728 CB TYR A 91	1.00 27.34
ATOM	700	1.00 42.02
ATOM		48.066 -3.039 99.194 1.00 24.28
		48.374 -1.829 99.822 1.00 21 55
MOTA	770	47.970 -0.609 99.275 1.00 24.69
MOTA		47.341 -2.997 98.002 1.00 24.86
MOTA	73.3 CE2 TYR A 91	46.931 -1.786 97.447 1.00 29 92
MOTA	734 CZ TYR A 91	47.250 -0.597 98.086 1.00 29.04
MOTA	735 OH TYR A 91	46.861 0.593 97.516 1.00 29.51
ATOM	736 C TYR A 91	47.452 -3.777 101.998 1.00 27.52
MOTA	737 O TYRA 91	
MOTA	738 N GLUA 92	16 100 27.20
MOTA	739 CA. GLU A 92	15 222
MOTA	740 CB GLU A 92	15 15 15 15 15 15 15 15 15 15 15 15 15 1
ATOM	741 CG GLU A 92	1.00 26.37
MOTA	742 CD GLU A 92	1 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2
ATOM	743 OE1 GLU A 92	44.178 1.395 102.396 1.00 37.40
ATOM	744 OE2 GLU A 92	42.999 1.293 102.794 1.00 31.22
ATOM		44.527 2.209 101.516 1.00 40.54
		45.770 -2.343 104.259 1.00 29.20
ATOM		46.389 -2.015 105.268 1.00 21.86
ATOM	747 N . ASN A 93	44.687 -3.117 104.286 1.00 26 51
ATOM	748 CA ASN A 93	44.109 ~3.613 105.527 1.00 24.02
ATOM	749 CB ASN A 93	42.727 -2.988 105.690 1.00 24.51
MOTA	750 CG ASN A 93	42.738 -1.488 105.405 1.00 28 61
ATOM	751 OD1 ASN A 93	43.428 -0.727 106.079 1.00 25.30
MOTA	752 ND2 ASN A 93	41.987 -1.063 104.393 1.00 20.45
ATOM	753 C ASN A 93	43.999 -5.132 105.407 1.00 24.79
ATOM	754 O ASNA 93	42.905 -5.680 105.291 1.00 21.89
ATOM	755 N PRO A 94	45.142 -5.828 105.429 1.00 24.60
MOTA	756 CD PRO A 94	46.493 -5.246 105.540 1.00 22.93
MOTA	757 CA PRO A 94	45.241 -7.285 105.312 1.00 27.23
ATOM	758 CE PRO A 94	46.730 -7.488 105.093 1.00 25.46
ATOM	759 CG PRO A 94	
ATOM	760 C PRO A 94	
ATOM	761 O PRO A 94	44 424
MOTA	762 N VAL A 95	44 606
ATOM	763 CA VAL A 95	14 200
ATOM	764 CB VAL A 95	47 070 ** ***
ATOM	765 CG1 VAL A 95	43 745 44
ATOM	766 CG2 VAL A 95	
ATOM	767 C VAL A 95	
ATOM	768 0 VAL A 95	46 637 44 444
ATOM	769 N SER A 96	45 664
ATOM	770 CA SER A 96	46 305
ATOM	771 CB SER A 96	
ATOM	772 OG SER A 96	46 750
ATOM	773 C SER A 96	
ATOM	774 C SER A 96	45.681 -10.804 111.854 1.00 32.10
ATOM	775 N TYR A 97	44 458 -10.839 111.950 1.00 37.91
ATOM		46.484 -10.795 112.913 1.00 32.57
		45.914 -10.801 114.248 1.00 34.95
ATOM		46.685 -11.735 115.182 1.00 35.47
ATOM		46.492 -13.187 114.817 1.00 40.65
ATOM	779 CD1 TYR A 97	47.319 -13.812 113.882 1.00 40.63
ATOM	780 CE1 TYR A 97	47.083 -15.121 113.475 1.00 42.16
ATOM	781 CD2 TYR A 97	45.421 -13.910 115.338 1.00 38.82
ATOM	782 CE2 TYR A 97	45.175 -15.219 114.936 1 00 42 82
ATOM	783 CE TYR A 97	46.010 -15.816 114.005 1.00 42.56
ATOM	784 CH TYR A 97	45.772 -17.105 113.601 1.00 46.03
MOTA	785 C TYR A 97	45.862 -9.394 114.813 1.00 37.56
ATCM	786 O TYRA 97	45.601 -9.195 115.998 1.00 39.06
ATOM ·	787 N ALA A 98	46.115 -8.418 113.948 1.00 31.96
ATOM	788 CA ALA A 98	46.048 -7.024 114.341 1.00 30.43
ATOM	789 CE ALA A 98	47.105 -6.211 113.600 1.00 29.64
ATOM	790 C ALA A .98	44.658 -6.533 113.962 1 00 30 35
ATOM	791 O ALA A 98	44.099 -5.655 114.612 1.00 31.82
ATOM	792 N MET A 99	44.094 -7.130 112.915 1.00 30.40
	 -	- 7.130 114.313 1.00 30.40

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793
 ATOM
                CA
                                     42.788
42.370
                    MET A
                           .99
                                              -6.730 112.420
                                                               1.00 27.54
 ATOM
           794
                CB
                    MET A
                            99
                                              -7.620 111.238
                                                                1.00 30.55
           795
 ATOM
                    MET A
                CG
                            99
                                              -9.071 111.565
                                     42.052
                                                                1.00 31.59
          796
 ATOM
                SD
                    MET A
                             99
                                     41.902
                                            -10.077 110.053
                                                                1.00 30.13
 ATOM
          797
                CE
                    MET A
                            99
                                     40.770
                                              -9.086 109.085
                                                               1.00 28.02
 ATOM -
          798
                C
                    MET A
                            99
                                     41.703
                                              -6.696 113.490
                                                               1.00 28.02
 ATOM
          799
                                     40.818
                0
                    MET A
                            99
                                              -5.842 113.446
                                                               1.00 24.53
 ATOM
          800
                    PHE A 100
                N
                                     41.752
                                              -7.614 114.449
                                                               1.00 26.07
 MOTA
          801
                CA
                                     40.759
                    PHE A 100
                                              -7.583 115.516
                                                               1.00 30.47
 ATOM
          802
                CB
                    PHE A 100
                                     39.738
                                              -8.718 115.404
                                                               1.00 30.29
 MOTA
          803
                CG
                    PHE A 100
                                              -8.657 116.475
                                     38.693
                                                               1.00 29.35
 MOTA
          804
                CD1
                    PHE A. 100
                                     37.722
                                             -7.662 116.455
                                                               1.00 27.01
 MOTA
          805
                    PHE A 100
                CD2
                                     38.756
                                             -9.506 117.575-
                                                               1.00 30.68
 ATOM
          806
                CE1
                    PHE A 100
                                             -7.507 117.519
                                     36.834
                                                               1.00 31.41
 MOTA
          807
                CE2
                    PHE A 100
                                     37.873
                                             -9.356 118.644
                                                               1.00 28.39
          808
                    PHE A 100
 MOTA
               CZ
                                     36.913
                                             -8.355
                                                     118.618
                                                               1.00 24.06
 MOTA
          809
               C
                    PHE A 100
                                     41.345
                                             -7.616 116.922
                                                               1.00 29.67
 MOTA
          810
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                    PHE A 100
                                     41.028
                                             -6.751 117.740
                                                               1.00 29.67
 ATOM
          811
               N
                    THR A 101
                                                               1.00 31.30
                                     42.181
                                             -8.610 117.222
 ATOM
          812
               CA
                    THR A 101
                                     42.770
                                             -8.701 118.562
                                                               1.00 31.37
 MOTA
          813
               CB
                    THR A 101
                                     43.610
                                             -9.977 118.732
                                                               1.00 31.63
 ATOM
          814
               0G1
                    THR A 101
                                     42.777 -11.119
                                                     118.532
                                                               1.00 31.64
 MOTA
          815
               CG<sub>2</sub>
                   THR A 101
                                    44.197 -10.045
                                                    120.137
                                                               1.00 27.38
 MOTA
          816
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                    THR A 101
                                    43.647
                                             -7.493 118.884
                                                               1.00 31.66
 ATOM
          817
               0
                    THR A 101
                                    43.502
                                             -6.875
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 ATOM
         818
               N
                    GLY A 102
                                             -7.166 117.976
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                                                               1.00 27.40
 ATOM
         819
               CA
                   GLY A 102
                                             -6.018 118.193
                                    45.430
                                                               1.00 27.19
 ATOM
         820
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                    GLY A 102
                                    44.631
                                             -4.728 118.266
                                                               1.00 27.26
                    GLY A 102
 MOTA
         821
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                                    44.785
                                             -3.940 119.201
                                                              1.00 27.68
         822
 MOTA
               N
                   SER A 103
                                    43.767
                                             -4.515 117.279
                                                              1.00 30.52
 ATOM
         823
               CA
                   SER A 103
                                            -3.314 117.216
                                    42.941
                                                               1.00
                                                                    31.91
 ATOM
         824
               CB
                   SER A 103
                                    42.085
                                            -3.334 115.949
                                                              1.00 34.63
 ATOM
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               OG
                   SER A 103
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                                             -3.265 114.791
                                                              1.00 35.94
 ATOM
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               C
                   SER A 103
                                    42.046
                                            -3.163 118.441
                                                              1.00
 ATOM
         827
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                                    41.891
                                            -2.065 118.984 1.00
ATOM
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              N
                   SER A 104
                                    41.455
                                            -4.270
                                                    118.871
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ATOM
         829
              CA
                   SER A 104
                                    40.584
                                            -4.251 120.038
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                                                                   30.22
ATOM
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              CB
                   SER A 104
                                    39.978
                                            -5.633 120.265
                                                              1.00 23.88
MOTA
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              OG
                   SER A 104
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                                            -5.595
                                                    121.358
                                                              1.00 36.91
ATOM
         832
              С
                   SER A 104
                                    41.367
                                            -3.841 121.282 - 1.00
                                                                   28.13
              O
ATOM
         833
                   SER A 104
                                    40.872
                                            -3.098 122.130
                                                              1.00 25.16
ATOM
         834
              N
                   LEU A 105
                                    42.594
                                            -4.336
                                                    121.386
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                   LEU A 105
ATOM
              CA
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MOTA
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ATOM
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                   LEU A 105
                                   45.461
                                            -5.176 123.754
                                                              1.00
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ATOM
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                                            -5.723 124.828
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              CD2
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MOTA
                  LEU A 105
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                                            -6.178 123.462
                                                              1.00 40.23
MOTA
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              С
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                                   43.834
                                            -2.552 122.511
                                                              1.00 32.09
ATOM
        841
              0
                  LEU A 105
                                   43.896
                                            -1.894 123.554
                                                              1.00 30.38
                  ALA A 106
        842
              N
MOTA
                                   44.081
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                                                              1.00 30.26
MOTA
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              CA
                  ALA A 106
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                                                              1.00
                                                                   28.31
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                  ALA A 106
ATOM
        844
              CB
                                            -0.386 119.738
                                                              1.00 23.88
                  ALA A 106
        845
              C
MOTA
                                   43.243
                                             0.268 121.434
                                                              1.00 26.04
ATOM
        846
              0
                  ALA A 106
                                   43.380
                                             1.376 121.952
                                                              1.00 20.63
                  THR A 107
THR A 107
              N
ATOM
        847
                                   42.058
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                                                              1.00 26.86
              CA
        848
MOTA
                                   40.841
                                             0.542 121.322
                                                              1.00 25.04
              CB
MOTA
        849
                  THR A 107
                                   39,706
                                             0.007 120.443
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              OG1
                  THR A 107
        850
MOTA
                                   40.111
                                             0.092 119.069
                                                              1.00 24.62
                  THR A 107
             CG2
MOTA
        851
                                   38.439
                                             0.824 120.629
                                                              1.00 19.80
                                             0.503 122.798
        852
             C
                  THR A 107
                                   40.450
ATOM
                                                              1.00 27.90
                  THR A 107
        853
             О
ATOM
                                   40.039
                                             1.515 123.361
                                                              1.00 29.04
                  GLY A 108
ATOM
        854
             Ν
                                   40.585
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                                                              1.00 24.01
        855
             CA
                  GLY A 108
MOTA
                                   40.256
                                            -0.767 124.832
                                                              1.00 24.86
        856
             C
                  GLY A 108
MOTA
                                   41.181
                                             0.155 125.603
                                                             1.00 23.86
                  GLY A 108
        857
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                                            0.790 126.572
ATOM
                                   40.771
                                                             1.00 26.97
                  SER A 109
        858
                                            0.236 125.158
ATCM
                                   42.434
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ATOM	859 CA SER A 109	43.421	1.090 125.807	1 00 00 05
ATOM	860 CB SER A 109	44.795	0.910 125.160	
ATOM	861 OG SER A 109	45.294	-0.393 125.402	
ATOM	862 C SER A 109	43.008	2.552 125.759	
ATOM	863 O SER A 109	43.323	3.312 126.672	1.00 21.13
ATOM	864 N THR A 110	42.311	2.949 124.698	1.00 23.17.
ATOM	865 CA THR A 110	41.841	4.327 124.583	1.00 20.83
ATOM	866 CB THR A 110	41.332		1.00 21.84
ATOM	867 OG1 THR A 110	42.452	4.648 123.161 4.769 122.276	1.00 24.33
ATOM	868 CG2 THR A 110	40.543	5.954 123.144	1.00 25.38
ATOM	869 C THR A 110	40.725	4.561 125.600	1.00 21.18
ATOM	870 O THR A 110	40.632	5.637 126.197	1.00 28.52
ATOM	871. N VAL A 111	39.882	3.558 125.809	1.00 28.27
ATOM .	872 CA VAL A 111	38.811	3.706 126.793	1.00 26.88
ATOM	873 CB VAL A 111	37.820	2.519 126.742	1.00 30.04
MOTA	874 CG1 VAL A 111	36.737	2.693 127.802	1.00 29.94
ATOM	875 CG2 VAL A 111	37.193	2.431 125.355	1.00 27.07
ATOM	876 C VAL A 111	39.440	3.797 128.187	1.00 25.26
ATOM	877 O VAL A 111	38.968	4.539 129.039	1.00 28.10
ATOM	878 N GLN A 112	40.521	3.056 128.415	1.00 26.06
ATOM	879 CA GLN A 112	41.188	3.097 129.711	1.00 23.92
ATOM	880 CB GLN A 112	42.268	2.020 129.804	1.00 30.27
ATOM	881 CG GLN A 112	41.777	0.629 129.481	1.00 28.61
ATOM	882 CD GLN A 112	42.883	-0.397 129.564	1.00 28.90
MOTA	883 OE1 GLN A 112	43.344	-0.740 130.653	1.00 28.60 1.00 29.68
ATOM	884 NE2 GLN A 112	43.333	-0.880 128.409	1.00 29.68
ATOM	885 C GLN A 112	41.834	4.461 129.931	1.00 22.13
MOTA	886 O GLN A 112	41.791	5.006 131.035	1.00 29.99
ATOM	887 N ALA A 113	42.453	5.004 128.885	1.00 28.64
ATOM	888 CA ALA A 113	43.083	6.315 129.001	1.00 26.62
MOTA	889 CB ALA A 113	43.693	6.732 127.684	1.00 23.49
ATOM :	890 C ALA A 113	42.005	7.307 129.407	1.00 24.63
ATOM	891 O ALA A 113	42.232	8.183 130.240	1.00 26.38
ATOM	892 N ILE A 114	40.824	7.163 128.822	1.00 25.26
ATOM	893 CA ILE A 114	39.728	8.063 129.145	1.00 27.05
MOTA	894 CB ILE A 114	38.554	7.887 128.156	1.00 26.93
ATOM	895 CG2 ILE A 114	37.387	8.770 128.576	1.00 25.86
MOTA	896 CG1 ILE A 114	39.008	8.259 126.739	1.00 28.38
MOTA	897 CD1 ILE A 114	37.938	8.105 125.669	1.00 28.64
ATOM	. 898 C ILE A 114	39.239	7.823 130.578	1.00 31.36
ATOM	899 O ILE A 114	38.898	8.770 131.291	1.00 24.56
ATOM	900 N GLU A 115	39.210	6.563 131.005	1.00 31.17
ATOM	901 CA GLU A 115	38.750	6.257 132.358	1.00 32.12
ATOM	902 CB GLU A 115	38.729	4.744 132.607	1.00 32.15
ATOM	903 CG GLU A 115	37.904	3.947 131.598	1.00 32.84
ATOM	904 CD GLU A 115 905 OE1 GLU A 115	37.875	2.459 131.912	1.00 34.12
ATOM		38.910	1.910 132.345	1.00 30.36
ATOM ATOM		36.826	1.827 131.699	1.00 31.38
ATOM	907 C GLU A 115 908 O GLU A 115	39.675	6.932 133.357	1.00 31.65
ATOM	909 N GLU A 116	39.224	7.446 134.383	1.00 29.25
ATOM	910 CA GLU A 116	40.970	6.933 133.053	1.00 31.50
ATOM	911 CB GLU A 116	41.942 43.367	7.564 133.934	1.00 32.34
ATOM	·912 CG GLU A 116		7.285 133.457	1.00 33.29
ATOM	913 CD GLU A 116	43.805 43.701	5.842 133.633	1.00 32.29
ATOM	914 OE1 GLU A 116	44.329	5.378 135.079 6.003 135.961	1.00 36.87
ATOM	915 OE2 GLU A 116	42.993	4.385 135.335	1.00 34.07
atom atom	916 C GLU A 116	41.702	9.067 134.006	1.00 35.00
ATOM	917 O GLU A 116	41.863	9.678 135.066	1.00 36.69
ATOM	918 N PHE A 117	41.317	9.661 132.881	1.00 34.39
ATOM	919 CA PHE A 117			1.00 31.19
ATOM	920 CB PHE A 117		11.509 131.444	1.00 28.43
ATOM	921 CG PHE A 117		12.908 131.381	1.00 30.14 1.00 35.78
ATOM	922 CD1 PHE A 117		14.006 131.501	1.00 33.78
RTOM	923 CD2 PHE A 117			1.00 33.39
MOTA	924 CE1 PHE A 117		15.299 131.466	1.00 30.65
·				1.00 20.03

ATOM	925		A 117	38.153	14.412	131.190	1.00 36.50)
ATOM	926		A 117	39.003	15.501	131.310	1.00 35.41	Ļ
ATOM	927		A 117	39.908		133.811	1.00 32.78	
ATOM ATOM	928 929		A 117 A 118	39.966		134.566	1.00 29.82	
ATOM	930		A 118 A 118	38.874 37.720	10.568	133.771 134.632	1.00 28.61	L
ATOM	931	CB LEU	A 118	36.621		134.632	1.00 32.00 1.00 29.19)
ATOM	932	CG LEU	A 118	36.098	9 830	132.820	1.00 29.19	,
ATOM	933	CD1 LEU	A·118	34.962		132.622	1.00 34.47	
ATOM	934	CD2 LEU	A 118	35.612		132.522	1.00 32.24	
MOTA	935		A 118	38.123	10.590	136.094	1.00 31.17	,
MOTA	936		A 118	37.576		136.964	1.00 28.32	:
ATOM	937		A 119	39.083		136.363	1.00 27.23	
MOTA MOTA	938 939		A 119 A 119	39.531 40.203		137.733	1.00 30.95	
ATOM	940		A 119	39.293	8.130	137.884 137.540	1.00 26.35	
ATOM	941		A 119	39.895		137.986	1.00 32.44 1.00 33.31	
ATOM	942		A 119 .	41.280		137.411	1.00 33.47	
ATOM	943		119	41.874	4.102	137.904	1.00 33.40	
ATOM	944		119	40.493	10.594	138.173	1.00 32.65	
ATOM	945		119	41.050	10.548	139.270	1.00 28.83	
ATOM	946 947	N GLY A	120	40.689		137.308	1.00 33.77	
ATOM ATOM	948	C GLY		41.571 43.035		137.652 137.340	1.00 33.84	
ATOM	949	O GLY		43.880	13.227		1.00 34.27 1.00 36.80	
ATOM		N ASN A		43.347	11.384		1.00 30.77	
MOTA	951	CA ASN A		44.731	11.122		1.00 31.73	
MOTA		CB ASN A		45.089		136.437	1.00 29.34	
MOTA		CG ASN A		44.856		137.851	1.00 35.83	
ATOM ATOM		OD1 ASN A ND2 ASN A		45.190 44.304		138.816	1.00 32.74	
ATOM		C ASN A		44.954		137,986 134.790	1.00 33.20 1.00 32.59	
ATOM		O ASN A		44.031		134.730	1.00 32.39	
ATOM	958	N VAL A		46.186		134.322	1.00 32.74	
MOTA		CA VAL A		46.540		132.946	1.00 33.59	
MOTA		CB VAL A		47.571		132.882	1.00 36.05	
MOTA		CG1 VAL A CG2 VAL A		47.884 47.029		131.438	1.00 37.58	
ATOM ATOM		C VAL A		47.029		133.602 132.352	1.00 37.19 1.00 34.47	
ATOM		O VAL A		48.053	9.801	132.939	1.00 31.28	
MOTA	965 1	N ALA A	123	46.646		131.196	1.00 28.06	
MOTA		CA ALA A		47.142	8.784	130.563	1.00 30.73	
ATOM		CB ALA A		46.133		130.727	1.00 32.69	
ATOM		C ALA A C ALA A		47.466 46.909		129.088	1.00 30.55	
MOTA MOTA		V PHE A		48.380		128.406 128.613	1.00 32.89 1.00 27.53	
ATOM		CA PHE A	124	48.807		127.229	1.00 27.33	
ATOM	972	CB PHE A		50.261		127.157	1.00 25.32	
ATOM		CG PHE A	124	50.903		125.793	1.00 27.84	
MOTA		D1 PHE A		50.179		124.629	1.00 24.77	
MOTA		D2 PHE A		52.266	8.266	125.686	1.00 21.79	
MOTA		E1 PHE A E2 PHE A		50.802		123.385	1.00 29.19	
ATOM ATOM		E2 PHE A		52.894 52.164		124.449 123.296	1.00 27.38 1.00 20.91	
ATOM	979 C			48.671		126.675	1.00 20.91	
ATOM	980 O			49.181		127.260	1.00 25.38	
ATOM	981 N			47.933		125.580	1.00 18.87	
MOTA		A ASN A		47.750	5.342	124.905	1.00 25.05	
MOTA		B ASN A		46.271		124.756	1.00,22.99	
ATOM		G ASN A D1 ASN A		46.073		123.856	1.00 24.08	
ATOM		D1 ASN A D2 ASN A		46.916 44.960		123.822 123.138	1.00 20.46	
ATOM ATOM	987 C			48.380		123.518	1.00 16.10 1.00 23.43	
ATOM	988 0			47.718	5.749		1.00 23.43	
ATOM	989 N		126	49.680	5.103	123.423	1.00 24.55	
MOTA	990 C	D PRO A	126	50.589	4.730	124.519	1.00 22.87	

* mo:	V 001 02 5			
ATO!		50.413	·5.130 122.16	1.00 22.39
ATO	M 992 CB PRO A 126	51.829	4.751 122.59	
ATO				
		51.564		
ATO		49.867	4.224 121.058	3 1.00 23.18
ATO	M 995 O PROA 126	50.173	4.436 119.893	1.00 20.12
ATO		49.058		
ATON				
		48.493		1.00 23.89
OTA		48.176	0.967 121.118	1.00 24.82
ATO 1	1 999 C ALA A 127	47.241		1.00 24.76
1OTA		46.806		
ATON				
		- 46.666		
ATOM		45.461	4.494 119.809	1.00 21.43
· ATOM	1 1003 C GLY A 128	45.732		1.00 23.55
ATOM	1 1004 O GLY A 128	46.875	5.695 118.291	
ATOM			5.095 116.291	
ATOM		44.680		
		44.822		1.00 24.99
ATOM		44.600	6.655 115.847	1.00 25.11
ATOM	1008 O GLY A 129	44.963	7.293 114.857	
ATOM		44.002		1.00 24.99
ATOM				
		43.729	4.825 114.481	
ATOM		43.360	3.361 114.744	1.00 22.77
MOTA	1012 CG MET A 130	44.455	2.661 115.563	1.00 26.30
ATOM		44.198		
ATOM			0.913 115.989	
		42.665	1.030 116:936	
ATOM		42.580	5.617 113.869	1.00 23.70
ATOM	1016 O MET A 130	41.421	5.199 113.901	1.00 26.28
ATOM	1017 N HIS A 131	42.926	6.766 113.294	1 00 20.20
ATOM	1018 CA HIS A 131	41.933	0.700 113.294	1.00 20.66
MOTA			7.687 112.775	1.00 20.99
		42.474	9.125 112.891	1.00 21.01
ATOM	1020 CG HIS A 131	43.699	9.391 112.069	1.00 28.30
MOTA	1021 CD2 HIS A 131	44.498	8.549 111.373	1.00 19.65
ATOM	1022 ND1 HIS A 131	44.246	10.649 111.917	
ATOM	1023 CE1 HIS A 131		10.045 111.91/	1.00 27.76
	1025 CDI 1115 A 151	45.328	10.567 111.163	1.00 20.48
MOTA	1024 NE2 HIS A 131	45.503	9.302 110.820	1.00 24.18
MOTA	1025 C HIS A 131	41.280	7.513 111.416	1.00 23.76
MOTA	1026 O HIS A 131	40.453	8.341 111.051	1.00 21.95
MOTA	1027 N HIS A 132	41.600	6.449 110.682	
ATOM	1028 CA HIS A 132			1.00 25.12
		41.006	6.257 109.354	1.00 23.32
MOTA	1029 CB HIS A 132	42.060	5.715 108.388	1.00 17.87
atom	1030 CG HIS A 132	43.148	6.689 108.072	1.00 24.79
MOTA	1031 CD2 HIS A 132	44.496	6.574 108.144	1.00 21.72
MOTA	1032 ND1 HIS A 132	42.896	7.944 107.556	
ATOM	1033 CE1 HIS A 132		7.544 107.556	1.00 13.58
		44.044	8.558 107.323	1.00 15.41
ATOM	1034 NE2 HIS A 132	45.028	7.748 107.668	1.00 15.27
ATOM	1035 C HIS A 132	39.752	5.386 109.208	1.00 23.38
ATOM	1036 O HIS A 132	38.947	5.615 108.304	2 00 24.70
ATOM	1037 N ALA A 133	39.587	4.388 110.070	
ATOM	1038 CA ALA A 133	38.453		1.00 23.34
			3.471 109.953	2 00 23.77
MOTA		38.515	2.417 111.053	1.00 27.49
ATOM	1040 C ALA A 133	37.093	4.145 109.966	1.00 23.02
MOTA	1041 O ALA A 133	36.878	5.117 110.691	1.00 25.98
ATOM	1042 N PHE A 134	36.179	3.633 109.148	
ATOM	1043 CA PHE A 134		3.633 109.148	1.00 18.90
	•	34.831	4.173 109.103	1.00 23.73
ATOM	1044 CB PHE A 134	34.317	4.296 107.663	1.00 24.29
MOTA	1045 CG PHE A 134	35.119	5.225 106.801	1.00 26.67
ATOM	1046 CD1 PHE A 134	36.025	4.724 105.867	
ATOM	1047 CD2 PHE A 134	34.975		1.00 28.69
			6.605 106.921	1.00 32.49
MOTA		36.775	5.582 105.063	1.00 28.65
ATOM	1049 CE2 PHE A 134	35.724	7.479 106.119	1.00 27.86
ATOM	1050 CZ PHE A 134	36.623	6.967 105.188	1.00 23.93
MOTA	1051 C PHE A 134	33.894	3.260 109.884	1.00 25.91
ATOM	1052 O PHE A 134	34.270		
			2.172 110.319	1.00 27.20
MOTA	1053 N LYS A 135	32.670	3.728 110.062	1.00 29.14
ATCM				
	1054 CA LYS A 135	31.638	2.984 110.765	1.00 35.26
ATOM	1054 CA LYS A 135 1055 CB LYS A 135	31.638 30.294		1.00 35.26
atom atom			2.984 110.765 3.628 110.429 2.779 110.667	1.00 35.26 1.00 35.86 1.00 46.26

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Figure 18-17

MOTA	1057	7 CD LYS A 135	27.834	3.542 110.211	1.00 47.72
MOTA	1058		26.610		1.00 53.65
ATOM	1059		26.788	1.549 109.167	1.00 53.27
ATOM	1060		31.617	1.490 110.414	1.00 35.62
ATOM	1061		31.609	0.635 111.301	
					1.00 32.58
ATOM-			31.629	1.180 109.122	1.00 35.97
MOTA	1063		31.555	-0.211 108.684	1.00 38.99
ATOM	1064		30.172	-0.474 108.083	1.00 38.87
MOTA	1065		29.146	-0.072 108.975	1.00 43.54
MOTA	1066		32.608	-0.616 107.660	1.00 37.84
MOTA	1067	' O SER A 136	32.350	-1.491 106.828	1.00 36.33
MOTA	1068	N ARG A 137	33.788	-0.008 107.705	1.00 33.23
MOTA	1069	CA ARG A 137	34.797	-0.368 106.724	1.00 30.89
MOTA	1070	CB ARG A 137	34.456	0.291 105.385	1.00 33.88
MOTA	1071	CG ARG A 137	35.009	-0.465 104.201	1.00 44.41
ATOM	1072	CD ARG A 137	34.809	0.261 102.880	1.00 46.27
MOTA	1073		35.091	-0.645 101.768	1.00 48.87
ATOM	1074		35.352	-0.261 100.526	1.00 48.64
ATOM	1075		35.372	1.029 100.220	1.00 51.82
ATOM	1076		35.592	-1.169 99.589	1.00 49.01
ATOM	1077		36.209	0.021 107.143	1.00 31.84
ATOM	1078	O ARG A 137	36.428	1.079 107.742	
	1079				1.00 30.36
ATOM	1080	N ALA A 138	37.166	-0.845 106.828	1.00 30.06
MOTA		CA ALA A 138	38.560	-0.588 107.158	1.00 32.24
ATOM	1081	CB ALA A 138	39.367		1.00 31.25
ATOM	1082	C ALA A 138	39.095	0.449 106.187	1.00 29.49
MOTA	1083	O ALA A 138	38.612	0.551 105.063	1.00 26.11
MOTA	1084	N ASN A 139	40.099	1.206 106.615	1.00 29.54
ATOM	1085	CA ASN A 139	40.673	2.241 105.767	1.00 26.99
ATOM	1086	CB ASN A 139	. 39.685	3.415 105.662	1.00 24.10
MOTA	1087	CG ASN A 139	40.209	4.556 104.811	1.00 28.02
ATOM	1088	OD1 ASN A 139	40.729	4.334 103.727	1.00 26.90
ATOM	1089	ND2 ASN A 139	40.050	5.789,105.293	1.00 23.55
ATOM	1090	C ASN A 139	42.027	2.713 106.285	1.00 30.17
ATOM	1091	O ASN A 139	42.245	2.827 107.497	1.00 27.55
ATOM	1092	N GLY A 140	42.944	2.959 105.354	1.00 31.82
ATOM	1093	CA GLY A 140	44.277	3.428 105.702	1.00 24.90
ATOM	1094	C GLY A 140	45.000	2.696 106.816	1:00 27.79
ATOM	1095	O GLY A 140	45.560	3.339 107.705	1.00 23.85
ATOM	1096	N PHE A 141	45.006	1.365 106.768	1.00 24.35
ATOM	1097	CA PHE A 141	45.679	0.538 107.783	1.00 24.53
ATOM	1098	CB PHE A 141	47.031	1.146 108.197	1.00 26.40
ATOM	1099	CG PHE A 141	47.997	1.366 107.062	1.00 30.31
ATOM	1100	CD1 PHE A 141	49.145	2.125 107.269	1.00 31.60
ATOM	1101	CD2 PHE A 141	47.781	0.811 105.802	1.00 29.44
ATOM	1102	CE1 PHE A 141	50.066	2.331 106.243	1.00 30.44
MOTA	1103.		48.694	1.008 104.770	1.00 27.91
ATOM	1104	CZ PHE A 141	49.840	1.771 104.991	1.00 29.38
ATOM	1105	C PHE A 141	44.846	0.387 109.056	1.00 23.53
ATOM	1106	O PHE A 141	45.194	-0.399 109.941	1.00 23.09
ATOM	1107		43.760		
ATOM	1108	CA CYS A 142	42.925	1.099 110.356	1.00 23.87
ATOM	1109	CB CYS A 142	42.472	2.516 110.723	1.00 22.51
ATOM	1110	SG CYS A 142	43.828	3.683 111.072	1.00 27.62
ATOM	1111	C CYS A 142	41.694	0.205 110.233	1.00 24.20
	1112				
ATOM		•	40.932	0.307 109.272	1.00 24.12
ATOM	1113	N TYR A 143	41.498	-0.663 111.219	1.00-23.84
ATOM	1114	CA TYR A 143	40.335	-1.546 111.236	1.00 26.07
MOTA	1115	CB TYR A 143	40.728	-2.958 111.680	1.00 27.89
ATOM	1116	CG TYR A 143	41.829	-3.582 110.855	1.00 27.30
ATOM	1117	CD1 TYR A 143	43.169	-3.329 111.137	1.00 25.76
ATOM	1118	CE1 TYR A 143	44.185	-3.875 110.346	1.00-25.77
MOTA	1119	CD2 TYR A 143	41.526	-4.394 109.762	1.00 25.87
ATOM '	1120	CE2 TYR A 143	42.531	-4.941 108.967	1.00 23.10
MOTA	1121	CZ TYR A 143	43.854	-4.679 109.262	1.00 22.93
MOTA	1122	OH TYR A 143	44.849	-5.217 108.476	1.00 20.64

MOTA	1123 C TYR A 143	39.281	-0.991 112.193	1.00 24.56
MCTA	1124 O TYR A 143	38.085		1.00 24.50
	-		-1.030 111.905	1.00 24.88
MOTA	1125 N ILE A 144	39.734		1.00 23.77
MOTA	1126 CA ILE A 144	38.833	0.102 114.335	1.00 27.11
ATOM	1127 CB ILE A 144	38.871		1.00 24.56
ATOM	1128 CG2 ILE A 144	37.941	-0.120 116.690	1.00 23.47
ATOM	1129 CG1 ILE A 144	38.430	-2.169 115.346	
	1130 CD1 ILE A 144			
ATOM		38.535	-3.113 116.539	
ATOM	1131 C ILE A 144	39.248	1.550 114.627	
MOTA	1132 O ILE A 144	40.428	1.843 114.800	
ATOM	1133 N ASNA 145	38.277	2.453 114.669	1.00 22.04
ATOM	1134 CA ASN A 145	38.555	3.866 114.920	
MOTA	1135 CB ASN A 145	37.559	4.732 114.133	1.00 18.87
ATOM	1136 CG ASN A 145	37.956	6.205 114.091	
-	1137 OD1 ASN A 145			1.00 22.21
ATOM		38.223	6.823 115.124	1.00 22.47
ATOM	1138 ND2 ASN A 145	37.978	6.776 112.892	1.00 23.78
ATOM	1139 C ASN A 145	38.417	4.141 116.418	1.00 22.63
ATOM	1140 O ASN A 145	37.338	4.535 116.880	1.00 22.45
ATOM	1141 N ASN A 146	39.495	3.941 117.178	1.00 16.63
MOTA	1142 CA ASN A 146	39.423	4.160 118.628	1.00 23.57
ATOM	1143 CB ASN A 146	40.708	3.678 119.320	
			4.500 119.320	1.00 19.80
MOTA		41.924	4.508 118.967	1.00 27.81
MOTA	1145 OD1 ASN A 146	42.299	5.421 119.704	1.00 19.55
ATOM	1146 ND2 ASN A 146	42.544	4.202 117.827	1.00 19.55
MOTA	1147 C ASN A 146	39.079	5.602 119.023	1.00 26.32
MOTA	1148 O ASN A 146	38.452	5.827 120.059	1.00 28.34
ATOM	1149 N PRO A 147	39.512	6.605 118.231	1.00 28.46
ATOM	1150 CD PRO A 147	40.383	6.637 117.042	1.00 27.18
ATOM	1151 CA PRO A 147	39.150		
			7.972 118.618	1.00 24.15
MOTA		39.859	8.815 117.558	1.00 25.13
MOTA	1153 CG PRO A 147	41.081	7.959 117.235	1.00 30.05
MOTA	1154 C PRO A 147	37.618	8.136 118.578	1.00 26.71
MOTA	1155 O PRO A 147	37.017	8.760 119.456	1.00 24.93
MCTA	1156 N ALA A 148	36.989	7:557 117.562	1.00 21.42
MOTA	1157 CA ALA A 148	35.536	7.633 117.416	1.00 21.03
ATOM	1158 CB ALA A 148	35.112	7.044 116.072	1.00 19.98
ATOM	1159 C ALA A 148	34.838	6.891 118.552	1.00 20.49
MOTA		33.822	7.344 119.067	1.00 21.44
ATOM	1161 N VAL A 149	35.381	5.739 118.928	1.00 19.20
MOTA	1162 CA VAL A 149	34.818	4.950 120.016	1.00 24.61
ATOM	1163 CE VAL A 149	35.570	3.608 120.181	1.00 25.96
ATOM	1164 CG1 VAL A 149	35.158	2.918 121.485	1.00 26.58
ATOM	1165 CG2 VAL A 149	35.262	2.704 118.995	1.00 25.67
ATOM	1166 C VAL A 149	34.947	5.752 121.304	1.00 23.56
ATOM	1167 O VAL A 149	33.990	5.887 122.064	1.00 22.52
ATOM	1168 N GLY A 150	36.143		
			6.287 121.536	1.00 24.65
ATOM	1169 CA GLY A 150	36.390	7.074 122.731	1.00 22.82
ATOM	1170 C GLY A 150	35.477	8.281 122.838	1.00 25.46
ATOM	1171 O GLY A 150	34.919	8.564 123.904	1.00 23.17
ATOM	1172 N ILE A 151	35.327	9.001 121.733	1.00 24.38
ATOM	1173 CA ILE A 151	34.481	10.180 121.716	1.00 22.85
ATOM	1174 CB ILE A 151	34.610	10.928 120.371	1.00 24.45
	1175 CG2 ILE A 151	33.598	12.077 120.306	
ATOM	1176 CG1 ILE A 151			1.00 24.71
ATOM		36.041	11.462 120.222	1.00 28.02
ATOM	1177 CD1 ILE A 151	36.354	12.056 118.854	1.00 27.10
ROTE	1178 C ILE A 151	33.018	9.806 121.987	1.00 28.19
ATOM	1179 O ILE A 151	32.337	10.482 122.763	1.00 26.37
ATOM	1180 N GLU A 152	32.532	8.734 121.364	1.00 26.32
ATOM	1181 CA GLU A 152	31.149	8.314 121.601	1.00 30.07
ATOM	1182 CB GLU A 152	30.758	7.161 120.672	1.00 29.37
	1183 CG GLU A 152	30.609	7.543 119.194	1.00 27.68
ATOM		29.455		
ATOM			8.504 118.946	1.00 31.82
ATOM	1185 OE1 GLU A 152	29.139	8.777 117.773	1.00 33.51
ATOM ·	1186 OE2 GLU A 152	28.862	9.009 119.918	1.00 34.73
ATOM	1187 C GLU A 152	31.009	7.879 123.055	1.00 28.00
ATOM	1188 O GLU A 152	29.980	8.096 123.683	1.00 31.23

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		•
ATO	M 1189 N TYR A 153	32.054 7.253 123 583 1 00 28 7
ATO	M 1190 CA TYR A 153	22 055
ATO	M 1191 CB TYR A 153	77 408
ATOI	M 1192 CG TYR A 153	23 (17
ATO	M 1193 CD1 TYR A 153	33.617 5.839 126.759 1.00 33.13
ATON	100	33.111 4.647 127.280 1.00 35.43
ATON	X	33.321 4.298 128.619 1.00 33 55
ATON	170	34.329 6.677 127.611 1.00 34 20
ATOM	x x 133	34.544 6.342 128.944 1.00 35.34
ATOM		34.041 5.154 129.444 1.00 37.50
	LIK H 133	34.260 4.835 130.767 1 00 30 10
ATOM	773	31.828 8.022 125.857 1.00 32.71
ATOM	21 110	31.026 7.988 126.787 1.00 29.14
ATOM		32.538 9.102 125.552 1.00 29.65
ATOM	11 134	2.00 29.05
ATOM		1.00 32.87
ATOM		31.46
MOTA	1205 CD1 LEU A 154	25.08
ATOM	1206 CD2 LEU A 154	2. 2. 2. 2. 2. 2. 2. 2. 2. 2. 2. 2. 2. 2
ATOM	1207 C LEU A 154	
MOTA	1208 O LEU A 154	20 455
ATOM	1209 N ARG A 155	1,00 32,38
MOTA	1210 CA ARG A 155	
ATOM	1211 CB ARG A 155	
ATOM	1212 CG ARG A 155	28.661 11.502 123.405 1.00 36.32
ATOM	1213 CD ARG A 155	29.581 12.253 122.460 1.00 43.15
ATOM	1214 NE ARG A 155	29.100 12.201 121.023 1.00 41.10
ATOM	1215 CZ ARG A 155	27.936 13.047 120.768 1.00 44.00
ATOM	1216 NH1 ARG A 155	27.331 13.140 119.583 1.00 54.07
ATOM	1217 NH2 ARG A 155	12.441 118.540 1.00 51.61
ATOM	1218 C ARG A 155	26.291 13.948 119.424 1.00 51.76
ATOM	1219 O ARG A 155	28.112 10.821 125.745 1.00 36.25
ATOM	1220 N LYS A 156	27.270 11.433 126.397 1.00 39.00
ATOM	1221 CA LYS A 156	28.213 9.496 125.765 1.00 36.48
ATOM	1222 CB LYS A 156	27.315 8.698 126.587 1.00 39.06
ATOM	1000	27.460 7.213 126.256 1.00 41.88
ATOM	1223 CG LYS A 156 1224 CD LYS A 156	26.6/2 6.816 125.020 1.00 51 15
ATOM	1225 CE LYS A 156	27.169 7.505 123.781 1.00 55.56
ATOM	1226 NZ LYS A 156	26.117 7.502 122.676 1.00 55 63
ATOM	1227 C LYS A 156	24.993 8.425 123.013 1.00 49 15
MOTA	.1228 O LYS A 156	27.527 8.932 128.076 1.00 39 91
ATOM	1229 N LYS A 157	26.636 8.658 128.876 1.00 37.01
MOTA		28.703 9.431 128.448 1.00 37.73
ATOM		28.985 9.725 129.847 1.00 36 52
ATOM		30.493 9.700 130.122 1.00 35 64
ATOM	,	31.094 8.308 130.174 1.00 35 44
ATOM		30.509 7.510 131.335 1.00 31.28
ATOM		31.077 6.106 131.388 1.00 31 48
ATOM		30.464 5.310 132,493 1.00 36 30
ATOM		28.423 11.097 130.197 1.00 38.12
ATOM		28.531 11.547 131.336 1.00 37 61
ATOM		27.842 11.768 129.205 1.00 36 27
		27.257 13.074 129.452 1.00 34 31
ATOM	1240 C GLY A 158	27.972 14.293 128.894 1 00 36 36
ATOM	1241 O GLY A 158	27.438 15.399 128.963 1 00 32 06
ATOM	1242 N PHE A 159	29.170 14.117 128.344 1 00 73 00
ATOM	1243 CA PHE A 159	29.892 15.260 127.796 1.00 30.29
MOTA	1244 CB PHE A 159	31.346 14.892 127.504 1.00 28.62
MOTA	1245 CG PHE A 159	32.137 14.555 128.730 1.00 28.80
ATOM	1246 CD1 PHE A 159	32.043 13.300 129.310 1.00 30.41
ATOM	1247 CD2 PHE A 159	32.951 15.513 129.327 1.00 29.37
ATOM	1248 CE1 PHE A 159	30 740 40 554 555
MOTA	1249 CE2 PHE A 159	33 661 15 000 130 400
MOTA	1250 CZ PHE A 159	1.00 31.10
MOTA	1251 C PHE A 159	1.00 32.32
	1252 O PHE A 159	70 700
	1253 N LYS A 160	20 180
	1254 CA LYS A 160	2.00 30.20
		28.550 17.766 125.254 1.00 33.98

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Figure 18-20

	•	
ATO	OM 1255 CB LYS A 160	27 200 20 200
ATO	M 1256 CG LYS A 160	27.390 18.653 125.719 1.00 36.87
ATO	X 100	26.273 17.914 126 419 1 00 20 46
	65 313 A 160	
ATO:	01 DIS A 100	25 500 20
ATO:	M 1259 NZ LYS A 160	25 024 1100 50.8
ATO	210 71 100	23.324 19.534 129.008 1 nn 45 76
	- 215 A 100	29.484 18.616 124 394 1:00 33
ATO		20 002 10
ATO	M 1262 N ARG A 161	20 700
ATOI	M 1263 CA ARG A .161	
ATO	ILIO A .101	31.003 19.626 174 108 1 00 00 on
	1210 A 101	
ATON	ING A 101	20.45
ATOM	M 1266 CD ARG A 161	20 705
· ATOM	1 1267 NE ARG A 161	23.341 124.966 1 00 39 01
	12/G N 101	31.158 23.503 126 341 1 00 43 76
ATOM		20 200 00 00 00 00 00 00 00 00 00 00 00
ATOM	1 1269 NH1 ARG A 161	20 12
ATOM	1 1270 NH2 ARG A 161	23.11/ 22.985 127.274 1.00 45 05
		30.893 23.518 128 627 1 00 42 74
ATOM	- 1210 H TOI	
ATOM	1272 O ARG A 161	77 700 1- 1-1-1-1-1 4.00 32./1
ATOM	1 1273 N ILE A 162	33.732 19.090 125.092 1.00 20 52
		33.257 18.087 123.149 1 00 22 50
ATOM	TID A 102	
ATOM	1275 CB ILE A 162	34 145 17 18 18 18 18 18 18 18 18 18 18 18 18 18
ATOM	1276 CG2 ILE A 162	
ATOM	102 H 102	33.40/ 14.976 122.898 1 00 24 05
	A 102	
ATOM	1278 CD1 ILE A 162	22 564 255 250 30.25
ATOM	1279 C ILE A 162	
ATOM	11 102	33.333 1/.816 121.886 1 nn 26 46
ATOM		· 34.876 17.973 120.762 1.00 27.88
	1281 N LEU A 163	
MOTA	1282 CA LEU A 163	22 505
ATOM	1283 CB LEU A 163	20 20 40 40 40 40 40 40 40 40 40
ATOM	1284 CG LEU A 163	30.304 19.729 121.681 1.00 26 25
		19 676 20 120 100
ATOM	1285 CD1 LEU A 163	70 210
ATOM	1286 CD2 LEU A 163	
ATOM	1287 C LEU A 163	
ATOM		30.34/ 1/.416 120.792 1.00 27 00
MOTA	1289 N TYR A 164	#.00 Z3.Z3
ATOM	1290 CA TYR A 164	20 7.5
ATOM	1291 CB TYR A 164	33.747 16.241 119.010 1.00 26 07
ATOM		39.021 15.181 118.179 1.00 23.38
	1292 CG TYR A 164	
MOTA	1293 CD1 TYR A 164	
ATOM	1294 CE1 TYR A 164	42 420 22.49
ATOM		41.419 12.224 117.794 1.00 22.90
	1111	40.202 14.142 116.194 1.00 18.74
MOTA	1296 CE2 TYR A 164	
ATOM	1297 CZ TYR A 164	41 663
ATOM	1298 OH TYR A 164	40 505
ATOM		42.506 11.296 115.878 7 00 19 41
		40.798 16.923 118 138 1 00 21 67
MOTA	1300 O TYR A 164	
ATOM	1301 N ILE A 165	40 000 19.73
MOTA	1302 CA ILE A 165	42 140 25.61
ATOM		43.149 17.462 117.804 1.00 24 43
		43.963 18.396 118.717 1.00 26.41
MOTA	1304 CG2 ILE A 165	
ATOM	1305 CG1 ILE A 165	43 035 10 100 141.337 1.00 19.36
MOTA	1306 CD1 ILE A 165	
ATOM		43.083 20.402 120.299 1.00 25 05
ATOM	1308 O ILE A 165	14 570
ATOM	1309 N ASP A 166	
ATOM		
		45.022 15.386 115.228 1.00-27 11
ATOM	1311 CB ASP A 166	
ATOM	1312 CG ASP A 166	44 600 12.703 114.13/ 1.00 28.56
MOTA	1313 OD1 ASP A 166	46 021
ATOM		45.831 13.456 113.068 1.00 30.37
		43.995 12.437 113 717 1 00 22 27
	1315 C ASP A 166	46 330
ATOM	1316 O ASP A 166	46 705
	1317 N LEU A 167	42 462
	. 220 11 207	4/-432 15.597 115.227 1 00 23 43
		48.738 16.068 114.722 1.00 24.67
	1319 CB LEU A 167	10 (00 = 1.00 24.6/
TOM	1320 CG LEU A 167	10 11 10 10 10 10 10 10 10 10 10 10 10 1
	-C CLU A TO!	49.143 17.444 116.858 1.00 26.62

	7771	ont			
MOTA			50.249	17.845 117.821	1.00 25.88
ATOM	1322	CD2 LEU A 167	48.658	18.668 116.092	
ATOM	1323	C LEU A 167	49.405		
ATOM	1324	O LEU A 167			
			50.504		
MOTA	1325	N ASP A 168	48.736	13.977 113.488	1.00 24.69
ATOM-	1326	CA ASP A 168	49.244	12.975 112.555	
ATOM	1327	CB ASP A 168	48.209		
	1328				
ATOM		CG ASP A 168	48.722		
MOTA	1329	C ASP A 168	49.423	13.686 111.209	1.00 24.17
ATOM	1330	O _ASP A 168	48.629		
ATOM	1331	OD1 ASP A 168	49.085		
				- · · · · ·	
MOTA	1332	OD2 ASP A 168	48.777		
MOTA	1333	N ALA A 169	50.448	13.312 110.446	1.00 21.29
ATOM	1334	CA ALA A 169	50.693	13.927 109.140	
ATOM	1335	CB ALA A 169	52.068	13.498 108.601	
	1336				1.00 21.17
MOTA		C ALA A 169	49.612	13.636 108.093	1.00 26.57
ATOM	1337	O ALA A 169	49.641	14.204 107.000	1.00 26.90
ATOM	1338	N HIS A 170	48.673	12.746 108.406	1.00 21.63
MOTA	1339	CA HIS A 170	47.592		
				12.445 107.468	1.00 24.79
ATOM	1340	C HIS A 170	46.243	12.867 108.045	1.00 20.98
ATOM	1341	O HIS A 170	46.044	12.849 109.255	1.00 24.12
ATOM	1342	CB HIS A 170	47.550	10.950 107.131	1.00 23.17
MOTA	1343	CG HIS A 170	48.830		
					1.00 30.28
MOTA	1344	ND1 HIS A 170	49.842	9.982 107.385	1.00 31.00
MOTA	1345	CE1 HIS A 170	50.825	9.634 106.577	1.00 24.33
MOTA	1346	CD2 HIS A 170	49.224	10.329 105.273	1.00 22.88
ATOM	1347	NE2 HIS A 170	50.502	9.828 105.285	1.00 21.89
ATOM	1348	N HIS A 171			
			45.317	13.231 107.171	1.00 21.14
ATOM	1349	CA HIS A 171	43.993	13.661 107.591	1.00 25.57
ATOM	1350	CB HIS A 171	43.234	14.242 106.404	1.00 22.47
ATOM	1351	CG HIS A 171	41.857	14.719 106.746	1.00 29.75
ATOM	1352	CD2 HIS A 171	41.433	15.648 107.634	1.00 25.75
					1.00 25.58
MOTA	1353	ND1 HIS A 171	40.721	14.201 106.160	1.00 28.90
ATOM	1354	CE1 HIS A 171 .	39.656	14.787 106.676	1.00 25.35
MOTA	1355	NE2 HIS A 171	40.060	15.669 107.573	1.00 32.40
ATOM	1356	C HIS A 171	43.169	12.533 108.204	1.00 29.61
	1357				
MOTA		O HIS A 171	43.169	11.411 107.698	1.00 27.62
MOTA	1358	N CYS A 172	42.461	12.852 109.286	1.00 26.52
ATOM	1359	CA CYS A 172	41.610	11.897 109.987	1.00 24.82
MOTA	1360	CB CYS A 172	41.460	12.322 111.456	1.00 29.47
ATOM	1361	SG CYS A 172			
			40.959	14.065 111.717	1.00 25.69
ATOM	1362	C CYS A 172	40.237	11.797 109.314	1.00 28.21
ATOM	1363	O CYS A 172	39.211	12.131 109.914	1.00 26.78
ATOM	1364	N ASP A 173	40.213	11.332 108.066	1.00 22.05
ATOM	1365	CA ASP A 173	38.949	11.217 107.350	
	1366				1.00 27.39
MOTA			39.167	10.646 105.931	1.00 30.47
MOTA	1367	CG ASP A 173	39.824 ⁻	9.264 105.922	1.00 29.77
ATOM	1368	OD1 ASP A 173	39.886	8,658 104,830	1.00 21.14
ATOM	1369	OD2 ASP A 173	40.288	8.787 106.978	1.00 30.04
		C ASP A 173	30.200		
ATOM			37.895	10.400 108.105	1.00 27.86
ATOM	1371	O ASP A 173	36.720	10.762 108.120	1.00 23.47
ATOM	1372	N GLY A 174	38.309	.9.315 108.753	1.00 25.84
ATOM	1373	CA GLY A 174	37.344	8.513 109.490	1.00 28.49
		C CIV 3 174			1.00 28.49
ATOM		C GLY A 174	36.694	9.296 110.619	1.00 26.14
ATOM	1375	O GLY A 174	35.475	9.287 110.780	1.00 21.39
ATOM	1376	N VAL A 175	37.510	9.984 111.409	1.00 27.24
ATOM		CA VAL A 175	36.995	10.773 112.523	
					1.00 25.53
ATOM		CB VAL A 175	. 38.137	11.299 113.401	1.00 30.54
ATOM	1379	CG1 VAL A 175	37.565	12.105 114.566	1.00 28.02
ATOM		CG2 VAL A 175	38.973	10.129 113.911	1.00 21.30
		VAL A 175	36.163	11.955 112.035	
MOTA					1.00 25.01
ATCM		VAL A 175	35.130	12.282 112.623	1.00 21.60
ATOM	1383	N GLN A 176	36.601	12.594 110.957	1.00 25.43
MOTA		CA GLN A 176	35.854	13.730 110.426	1.00 26.12
		CB GLN A 176	36.554	14.336 109.205	1.00 24.71
ATOM				14.330 207.403	
ATOM	1386	G GLN A 176	35.682	15.349 108.469	1.00 26.68

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Figure 18-22

	1OTA			36.38	5 16.002	107.30	6 1.00 29	0 5/
	ATO I	4 1388 OF	El GLN A 176	37.38		107.48	1.00 26	7.J9
	ATON	1 1389 NE		35.87		106.09		
	ATOM	1 1390 C	GLN A 176	34.44		110.029		/.58
	ATOM		GLN A 176	33.48		110.023		/.63
	ATOM		GLU A 177		1 14.021	110.319		5.93
	ATOM		CIU A 177	34.33		109.362		2.21
				33.02		108.915		2.72
	ATOM			33.18		108.053		1.20
	ATOM			31.90	5 10.069	107.329	1.00 39	.40
	ATOM			32.06	0 8.819	106.497	1.00 41	
	ATOM	1397 OE	1 GLU A 177	32.05	6 7.712	107.075	1.00 45	
	ATOM	1398 OE	2 GLU A 177	32.20		105.264	1.00 42	35
	ATOM		GLU A 177	32.12		110.099		
	ATOM	1400 o	GLU A 177	30.94		110.093		. 54
	ATOM	1401 N	ALA A 178	32.70	7 10 750	111.114		
	MOTA		ALA A 178		10.750	111.114		.03
	ATOM		ALA A 178	31.97	10.365	112.303		
	ATOM	•		32.905	9.658	113.289		.49
			ALA A 178	31.261	11.519	113.003		
	MOTA		ALA A 178	30.145	11.355	113.493	1.00 28	.64
	ATOM	1406 N	PHE A 179	31.888		113.055	1.00 29	.27
	ATOM	1407 CA	PHE A 179	31.256	13.801	113.751	1.00 30	.49
	ATOM	1408 CB	PHE A 179	32.071	14.128	115.001	1.00 24	44
	MOTA	1409 CG	PHE A 179	32.469	12.909	115.781	1.00 24	43
	MOTA	1410 CD1	PHE A 179	33.749	12.375	115.657	1.00 25	
	. ATOM	1411 CD2	PHE A 179	31.536	12.233	116.563	1.00 23	
	ATOM	1412 CE1	PHE A 179	34.103	11 184	116 203	1.00 23	.09
	ATOM	1413 CE2	PHE A 179	31.881	11 038	117.204	1 00 26	. 50
	ATOM	1414 CZ	PHE A 179	33.170	10 515	117.204	1.00 26	.38
	ATOM	1415 C	PHE A 179.	31.079			1.00 20	.30
	ATOM	1416 0	PHE A 179	31.006			1.00 31.	.00
	ATOM	1417 N	TYR A 180	30.980		113.399	1.00 31.	. 32
	ATOM	1418 CA	TYR A 180			111.584	1.00 31.	. 68
	ATOM	1419 CB	TYR A 180	30.829		110.646	1.00 32.	
	MOTA	1420 CG		30.931	15.378		1.00 35.	
			TYR A 180	31.331	16:406		1.00 36.	. 27
	ATOM	1421 CD1	TYR A 180	30.427	16.846	107.204	1.00 37.	31
	ATOM	1422 CE1	TYR A 180	30.801	17.791		1.00 34.	81
	MOTA	1423 CD2	TYR A 180	32.624	16.937		1.00 36.	32
	MOTA		TYR A 180	33.007	17.879 1	L07.203	1.00 37.	83
	ATOM	1425 CZ	TYR A 180	32.088	18.304 1	06.250	1.00 36.	
	ATOM	1426 ОН	TYR A 180	32.446	19.255 1		1.00 28.	04
	MOTA	1427 C	TYR A 180	29.518	16.696 1	10.825	1.00 30.	9/
	ATOM	1428 O	TYR A 180	29.459	17.894 1	10.560	1.00 30.	
	ATOM	1429 N	ASP A 181	28.473	16.026 1	11.299	1.00 31.	
	MOTA	1430 CA	ASP A 181	27.180	16.691 1	11 444	1.00 37.	
	ATOM	1431 CB	ASP A 181	26.086	15.833 1	10 807	1.00 37.	
	ATOM		ASP A 181	25.645	14.689 1	11 705	1.00 37.	00
	ATOM	1433 OD1 .		26.505	13.963 1	12 222		
	ATOM		ASP A 181	24.425	14.504 1	11 021	1.00 43.	45
•	ATOM		ASP A 181	26.754	17.044 1	12 056	1.00 46.	26
	ATOM		ASP A 181	25.571	17.286 1	12.000	1.00 36.	81
	ATOM				17.200 1	13.109	1.00 33.9	91
	ATOM		THR A 182	27.003	17.000 1		1.00 40.	
	ATOM			27.327	17.412 1	15.184	1.00 38.2	
			THR A 182	27.433	16.201 1	16.133	1.00 37.9	99
	MOTA		THR A 182	27.013	16.595 1	17.448	1.00 35.6	54
	ATOM		THR A 182	28.869	15.684 1		1.00 35.6	
	ATOM		THR A 182	28.177	18.546 13	15.746	1.00 39.5	
	ATOM		THR A 182	29.365	18.673 13	15.433	1.00 40.0	
	ATOM		ASP A 183	27.557		16.582	1.00 37.0	
	ATOM	1445 CA A	ASP A 183	28.250	20.497 11		1.00 37.7	
	ATOM		SP A 183	27.313	21.706 11	17.228	1.00 37.7	
	ATCM		SP A 183	26.136	21.493 11	8 155	1.00 33.3	
	STOM		SP A 183	25.614	20.357 11			
	ATOM		SP A 183	25.720			1.00 34.9	
	ATCM ·	1450 C A	SP A 183	28.762	22.470 11		1.00 38.1	
	ATCM		SP A 183	29.337	20.161 11	_	1.00 35.2	
	ATOM	1452 N G	LN A 184	28.562	21.015 11		1.00 35.1	
	A L CIT	.ي ن. د. د	THE 12 TO 2	20.204	18.917 11	5.012	1.00 35.1	0

ATOM	1453	CA	GLN A	184	29.030	18.505	120.333	1.00	35.16
ATOM	1454	CB	GLN A		28.155	17.382	120.906	1.00	36.94
ATOM	1455	CG	GLN A	184	26.663		120.988		38.34
MOTA	1456	CD	GLN A	184	25.881		121.838		43.68
MOTA	1457	OE1	GLN A	184	26.027		121.696		35.48
ATOM	1458	NE2	GLN A		25.036		122.723		51.06
ATOM	1459	С	GLN A		30.479		120.253		36.32
MOTA	1460	0	GLN A		31.135		121.275		34.24
MOTA	1461	N	VAL A		30.976		119.028		34.51
ATOM	1462	CA	VAL A		32.348	17.443	118.804 118.259	1.00	35.11
MOTA	1463	CB	VAL A		32.393		118.003		23.80
MOTA	1464	CG1			33.834 31.731	15.567	119.242		26.00
ATOM	1465	CGZ	VAL A		33.053	18 354	117.803		33.11
ATOM	1466 1467	o	VAL A		32.545	18.593	116.714		27.73
ATOM ATOM	1468	N	PHE A		34.215		118.184		31.49
ATOM	1469	CA	PHE A		34.985		117.291		30.63
ATOM	1470	CB	PHE A		35.420		117.991	1.00	30.34
ATOM	1471	CG	PHE A		36.008	22.047	117.051	1.00	30.22
ATOM	1472	CD1	PHE A	186	35.265		116.656		32.23
ATOM	1473	CD2	PHE A	186	37.284	21.879	116.524		29.37
ATOM	1474	CE1	PHE A		35.785		115.748		27.87
ATOM	1475	CE2	PHE A		37.813		115.615		28.54
MOTA	1476	CZ	PHE A		37.064		115.227		30.80
MOTA	1477	C	PHE A		36.232		116.879 117.729		28.30
ATOM	1478	0	PHE A		36.952 36.478	18 877	115.574		32.00
ATOM	1479 1480	N CA	VAL A		37.645		115.060		29.70
MOTA MOTA	1481	CB	VAL A		37.252		114.019		30.03
ATOM	1482	CG1	VAL A		38.510	16.488	113.405	1.00	27.77
ATOM	1483	CG2			36.410		114.672		25.98
ATOM	1484	С	VAL A	187	38.604		114.392		31.03
ATOM	1485	0	VAL A	187	38.215		113.491		31.88
MOTA	1486	N	LEU A		39.850		114.857		24.88
ATOM	1487	CA	LEU A		40.899		114.304		26.92 27.04
ATOM	1488	CB	LEU A		41.468	20.959			25.15
ATOM	1489	CG	LEU A		42.823 42.686		114.963 113.648		18.53
ATOM	1490	CD1 CD2	LEU A		43.330		116.068		28.81
ATOM	1491 1492	CDZ	LEU A		42.022		113.815		31.22
ATOM ATOM	1493	ō	LEU A		42.579	18.333		1.00	25.83
ATOM	1494	N	SER A		42.369	19.230	112.540		30.53
ATOM	1495	CA	SER A	189	43.429	18.399	112.007		30.13
MOTA	1496	CB	SER A		42.821	17.249	111.199		33.41
MOTA	1497	OG	SER A		43.837		110.588		32.98 27.94
MOTA	1498	C	SER A		44.448	19.120	111.143		27.94
ATOM	1499	0	SER A		44.084	19.891	11 1.253 11 .423		24.80
ATOM	1500	N	LEU A		45.728 46.805	10.077	110.614		22.23
MOTA	1501 1502 -	CA CB	LEU A		47.955	20.000	111.459		23.69
MOTA	1503				47.733		112.522		28.92
MOTA. MOTA	1504	CD1	LEU A	190	49.070		112.740	1.00	23.01
ATOM	1505		LEU A		46.691	22.093	112.087	1.00	28.11
MOTA	1506	C	LEU A		47.300	18.210	109.872		22.69
ATOM	1507	0	LEU A		47.416	17.141	110.465		16.55
ATOM	1508	N	HIS A	191	47.599	18.353	108.587		19.22
ATOM	1509	CA	HIS A		48.046		107.804		23.28
ATOM	1510	CB	HIS A		46.870	16.242			15.58
ATOM	1511	CG	HIS A		45.591	16.915	107.256		24.16 17.71
MOŢĶ	1512	CD2	HIS A	191	45.034	17.124	106.038 108.176	1 00	23.76
ATOM	1513	NDI	HIS A	191	44.695 43.644	17.419			19.78
ATOM	1514	CEI.	HIS A	191	43.823	17.746	106.246		27.87
ATOM	1515 1516	NE2 C	HIS A	191	48.570	17.620	106.434		23.65
ATCM	1517	0	HIS A		48.419	18.761	106.017	1.00	23.89
ATOM ATOM	1518	N	GLN A	192	49.209	16.681	105.746	1.00	23.49
4		••			-				

					•				
ATOM .	1519	CA	GLN A	192		49.718	16.950	104.412	1.00 20.55
ATOM	1520	CB	GL _i N A			50.474	15.738	103.864	1.00 23.63
			GLN A			51.528	15.181	104.797	1.00 24.07
ATOM	1521	CG						104.293	1.00 26.90
MOTA	1522	CD	GLN A			52.110			
MOTA	1523	OE1	GLN A	192		52.986		103.421	1.00 20.21
ATOM	1524	NE2	GLN A	192		51.605		104.828	1.00 23.52
			GLN A			48.478	17.174	103.570	1.00 21.41
MOTA	1525	C						103.726	1.00 20.15
MOTA	1526	0	GLN A	192		47.478			
MOTA	1527	N	SER A	193		48.528		102.692	1.00 24.36
ATOM	1528	CA	SER A	193		47.397	18.448	101.821	1.00 23.98
			SER A			47.760	19.537	100.820	1.00 24.60
MOTA	1529	CB			•		19.660	99.861	1.00 25.83
ATOM	1530	OG	SER A			46.729	19.000		
MOTA	1531	С	SER A	193		46.985	17.200	101.045	1.00 23.74
ATOM	1532	0	SER A	.193		47.829	16.492	100.506	1.00 19.80
	1533	N·	PRO A			45.674	16.936	100.953	1.00 24.85
ATOM						44.561		101.507	1.00 25.08
MOTA	1534	CD	PRO. A						
MOTA	1535	CA	PRO A	194		45.151		100.235	1.00 29.25
ATOM	1536	CB	PRO A	194		43.641		100.444	1.00 30.51
	1537		PRO A	394		43.554	16.643	101.758	1.00 30.21
MOTA			PRO A			45.527	15.825	98.756	1.00 30.75
MOTA	1538	C					14.830	98.041	1.00 30.04
MOTA	1539	. 0	PRO A			45.420			
ATOM	1540	N	GLU A	195		45.967	16.991	98.298	1.00 26.28
ATOM	1541	CA	GLU A	195		46.343	17.127	96.898	1.00 31.11
	1542	CB	GLU A			46.738	18.570	96.571	1.00 29.52
ATOM						45.680	19.600	96.933	1.00 38.32
MOTA	1543	CG	GLU A					96.352	1.00 44.15
ATOM	1544	CD	GLU A			45.976	20.972		
ATOM	1545	OE1	GLU A	195		47.139	21.425	96.434	1.00 44.23
ATOM	1546	OE2	GLU A	195		45.037	21.605	95.825	1.00 45.06
		C	GLU A			47.499	16.193	96.552	1.00 30.81
ATOM						47.582	15.705	95.426	1.00 37.17
ATOM	1548	0	GLU A					97.515	1.00 25.01
ATOM	1549	N	TYR A	196		48.377	15.922		
ATOM	1550	CA	TYR A	196		49.517	15.053	97.242	1.00 23.43
ATOM	1551	CB	TYR A	196		50.810	15.881	97.223	1.00 26.67
	1552	CG	TYR A			51.255	16.424	98.572	1.00 26.78
MOTA						51.957	15.625		1.00 26.08
ATOM	1553		TYR A					100.734	1.00 26.77
ATOM	1554	CE1	TYR A	196		52.338			
ATOM	1555	CD2	TYR A	196	•	50.944	17.731	98.958	1.00 27.55
ATOM	1556	CE2	TYR A	196		51.320	18.226	100.216	1.00 25.95
	1557	CZ	TYR A			52.012	17.409	101.096	1.00 24.78
ATOM			TYR A			52.356		102.345	1.00 25.50
MOTA	1558	OH					13.906		1.00 27.05
MOTA	1559	C.	TYR A			49.670			
ATOM	1560	0	TYR A	196		50.585	13.088		
ATOM	1561	N	ALA A	197		48.785	13.822	99.214	1.00 22.10
	1562	CA	ALA A	197		48.928	12.760	100.199	1.00 24.90
MOTA			ALA A			49.627	13.307	101.437	1.00 27.83
ATOM .	1563	CB						100.608	1.00 26.20
ATOM	1554	С	ALA A			47.644			1.00 22.82
ATOM	1: ō 5	0	ALA A			46.553		100.484	
ATOM	1556	N	PHE A	198		47.795		101.102	1.00 31.74
ATOM	1567	CA	PHE A	198		46.663		101.580	1.00 28.74
		CB	PHE A			47.130	8.691	102.036	1.00 30.66
MOTA	1568						7 766	102.399	1.00 29.61
MOTA	1569	CG	PHE A			46.009	6 070	103 457	1.00 28.76
ATOM	1570	CD1	PHE A	198		45.496		101.463	
ATOM	1571	CD2	PHE A	198		45.426		103.657	1.00 28.43
	1572	CEI	PHE A	198		44.415	6.057	101.773	1.00 35.72
MOTA		CDJ	PHE A	100		44.340	7,004	103.970	1.00 34.62
MOTA	1573		FRE A	100				103.029	1.00 35.73
ATOM	1574	cz	PHE A	TAR		43.837	0.121	100.023	1 00 00 05
ATOM	1575	С	PHE A	198		46.121	10.814	102.802	1.00 28.95
	1576	ō	PHE A	198		46.892		103.596	1.00 25.72
ATOM			PRO A	199		44.792		102.941	1.00 28.27
ATOM	.1577	N					11 /00	104.099	1.00 33.97
ATOM	1578	CD	PRO A			44.100	11.477	102.002	1.00 32.80
ATOM	1579	CA	PRO A	199		43.813	10.364	102.008	
ATOM	1580	СВ	PRO A	199		42.550	10.312	102.858	1.00-29.84
		CG	PRO A			42.665	11.612	103.592	1.00 37.32
ATOM	1581		PRO A	100		43.773	11 476	100.965	1.00 35.02
ATOM	1582	C					10 631	101.280	1.00 48.84
MOTA	1583	0	PRO A	199		44.052			1 00 33 64
ATOM	1584	N	PHE A	200		43.441	11.156	99.734	1.00 33.64
W. O									

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ATOM	1585	CA PHE A 200	43.41	8 12.179 98.718	1.00 28.12
	1586				
MOTA					
ATOM	1587	CG PHE A 200	45.22	6 10.833 97.561	1.00 27.33
MOTA	1588	CD1 PHE A 200	45.23	9 9.510 97.995	1.00 29.79
ATOM	1589	CD2 PHE A 200			
MOTA	1590	CE1 PHE A 200			
MOTA	1591	CE2 PHE A 200	47.65	1 10.782 97.473	1.00 31.41
ATOM	1592	CZ PHE A 200	47.65	3 9.458 97.906	1.00 29.64
	1593	C PHE A 200			
ATOM					
MOTA	1594	O _ PHE A 200			
ATOM	1595	N GLU A 201	41.00	2 12.101 98.979	1.00 28.52
ATOM	1596	CA GLU A 201	39.61	4 12.534 98.806	1.00 35.04
ATOM	1597	CB GLU A 201			
MOTA	1598	CG GLU A 201			
ATOM	1599	CD GLU A 201			
ATOM	1600	OE1 GLU A 201	36.99	2 9.142 97.825	1.00 40.96
ATOM	1601	OE2 GLU A 201	38.77	2 7.937 98.298	1.00 44.17
	1602	C GLU A 201	39.07		
MOTA					
MOTA	1603	0 GLU A 201	. 38.08		
ATOM	1604	N LYS A 202	39.693	3 13.552 101.007	1.00 34.63 [.]
MOTA	1605	CA LYS A 202	39.229	9 14.460 102.030	1.00 34.09
ATOM	1606	CB LYS A 202	38.29		
ATOM	1607	CG LYS A 202	37.013		
MOTA	1608	CD LYS A 202	35.93	5 12.854 103.230	1.00 47.39
MOTA	1609	CE LYS A 202	34.628	3 12.663 102.469	1.00 47.74
MOTA	1610	NZ LYS A 202	33.504	1 12.290 103.378	1.00 53.56
ATOM	1611	C LYS A 202	40.382		
MOTA	1612	O LYS A 202	41.520		1.00 28.06
ATOM	1613	N GLY A 203	40.080		
MOTA	1614	CA GLY A 203	41.115	16.862 104.228	1.00 33.75
ATOM	1615	C GLY A 203	41.288	3 18.288 103.729	1.00 30.54
ATOM	1616	O GLY A 203	42.174		
ATOM	1617	N PHE A 204	40.458		
ATOM	1618	CA PHE A 204	40.557		
ATOM	1619	CB PHE A 204	39.863	3 20.217 100.901	1.00 31.41
ATOM	1620	CG PHE A 204	40.498	3 19.416 99.803	1.00 31.06
ATOM	1621	CD1 PHE A 204	40.169		
ATOM	1622	CD2 PHE A 204	41.431		
MOTA	1623	CE1 PHE A 204	40.761		
MOTA	1624	CE2 PHE A 204	42.033	19.267 97.931	1.00 36.08
MOTA	1625	CZ PHE A 204	41.697	17.928 97.751	1.00 36.54
ATOM	1626	C PHE A 204	39.967		1.00 37.30
			39.088		1.00 33.56
ATOM	1627	O PHE A 204			
MOTA	1628	N LEU A 205	40.451		1.00 38.52
ATOM	1629	CA LEU A 205	40.012	23.427 103.993	1.00 36.81
ATOM .	1630	CB LEU A 205	40.801	24.695 103.659	1.00 34.73
ATOM	1631	CG LEU A 205	40.496		1.00 40.98
	1632	CD1 LEU A 205	-40.690		1.00 39.87
MOTA					
ATOM	1633	CD2 LEU A 205	41.415		1.00 39.94
ATOM	1634	C LEU A 205	38.520		1.00 36.58
ATOM	1635	O LEU A 205	37.931	24.178 104.905	1.00 40.98
ATOM	1636	N GLU A 206	37.909		1.00 36.07
			36.486		1.00 36.30
MOTA	1637				
ATOM	1638	CB GLU A 206	36.107		1.00 39.98
ATOM	1639	CG GLU A 206	36.890		1.00 48.04
MOTA	1640	CD GLU A 206	38.307	23.980 99.868	1.00 51.87
	1641	OE1 GLU A 206	39.146		1.00 50.32
MOTA					1.00 56.69
ATOM	1642	OE2 GLU A 206	38.581		
ATOM	1643	C GLU A 206	35.572		1.00 33.85
ATOM	1644	O GLU A 206	34.433		1.00 26.22
ATOM	1645	N GLU A 207	36.07i		1.00 31.68
	1646	CA GLU A 207	35.297		1.00 31.65
ATCM					1.00 34.15
ATOM	1647	CB GLU A 207	36.000		
ATOM	1648	CG GLU A 207	36.044		1.00 33.80
ATOM	1649	CD GLU A 207	37.182		1.00 33.85
ATOM	1650	OE1 GLU A 207	37.487		1.00 33.22
AION		- .			

ATOM	1651 OE2 GLU A 207		37.76	0 17.688 101.916	1.00 35.48
MOTA			35.18		
MOTA			36.009		
MOTA			34.150	22.024 106.302	
MOTA			33.968		1.00 38.96
MOTA			33.737	7 24.134 107.529	
ATOM			33.717	7 24.762 108.914	
ATOM	1658 CG1 ILE A 208	•	34.841		1.00 40.34
ATOM			36.207		1.00 46.23
MOTA	1660 C ILE A 208		32.821		1.00 38.32
ATOM	1661 O ILE A 208	-	32.558		1.00 40.08
MOTA	1662 N GLY A 209		32.142		
MOTA	1663 CA GLY A 209		31.047		
ATOM	1664 C GLY A 209 1665 O GLY A 209		29.699		1.00 37.87
MOTA MOTA	1665 O GLY A 209 1666 N GLU A 210		29.579		1.00 40.56
ATOM	1667 CA GLU A 210		28.676	19.917 108.380	1.00 37.38
MOTA	1668 CB GLU A 210		27.337	20.118 107.831	1.00 42.34
ATOM	1669 CG GLU A 210		27.008 26.860		1.00 42.73
ATOM	1670 CD GLU A 210		26.633		1.00 47.38
ATOM	1671 OE1 GLU A 210		26.385		1.00 52.68
ATOM	1672 OE2 GLU A 210		26.711	16.810 105.226	1.00 50.59 1.00 53.78
MOTA	1673 C GLU A 210		26.287	20.114 108.938	1.00 42.90
ATOM	1674 O GLU A 210		26.516	19.577 110.022	1.00 45.94
ATOM	1675 N GLY A 211		25.130	20.702 108.654	1.00 43.16
MOTA	1676 CA GLY A 211		24.068	20.751 109.642	1.00 43.98
ATOM	1677 C GLY A 211		24.514	21.450 110.911	1.00 45.01
MOTA	1678 O GLY A 211		25.186	22.479 110.858	1.00 48.15
MOTA	1679 N LYS A 212		24.145	20.896 112.059	1.00 41.63
MOTA	1680 CA LYS A 212		24.528	21.495 113.328	1.00 45.07
MOTA	1681 CB LYS·A 212 1682 CG LYS A 212		23.913	20.715 114.490	1.00 46.59
MOTA MOTA	1682 CG LYS A 212 1683 CD LYS A 212		22.386	20.591 114.462	1.00 55.31
ATOM	1684 CE LYS A 212		21.651	21.945 114.481	1.00 57.42
ATOM	1685 NZ LYS A 212		21.749 21.051	22.696 113.151	1.00 59.71
ATOM	1686 C LYS A 212		26.046	24.017 113.178 21.513 113.469	1.00 57.43
ATOM	1687 0 LYS A 212		26.598	22.326 114.207	1.00 42.08 1.00 40.03
ATOM	1688 N GLY A 213		26.713	20.615 112.751	1.00 40.03
ATOM	1689 CA GLY A 213		28.163	20.538 112.817	1.00 40.11
ATOM	1690 C GLY A 213		28.888	21.519 111.916	1.00 38.25
ATOM	1691 O GLY A 213		30.122	21.575 111.913	1.00 34.70
ATOM	1692 N LYS A 214		28.131	22.295 111.143	1.00 37.31
MOTA	1693 CA LYS A 214		28.736	23.274 110.250	1.00 39.58
ATOM	1694 CB LYS A 214		27.656	24.017 109.463	1.00 44.69
ATOM	1695 CG LYS A 214		28.189	25.030 108.461	1.00 44.53
ATOM	1696 CD LY: A 214		27.047	25.704 107.720	1.00 47.71
ATOM ATOM	1697 CE LY. A 214 1698 NZ LY. A 214		27.553	26.759 106.754	1.00 52.94
MOTA	1699 C LYS A 214		28.453 29.547	26.183 105.717	1.00 57.45
ATOM	1700 O LYS A 214		29.002	24.259 111.085 24.963 111.933	1.00 40.16
ATOM	1701 N GLY A 215		30.851		1.00 37.92
ATOM	1702 CA GLY A 215		31.716	25.183 111.593	1.00 36.57 1.00 35.03
ATOM	1703 C GLY A 215		32.431	24.448 112.709	1.00 34.57
ATOM	1704 O GLY A 215		33.216	25.039 113.454	1.00 33.76
MOTA	1705 N TYR A 216		32.168	23.153 112.837	1.00 34.61
ATOM	1706 CA TYR A 216		32.816	22.378 113.885	1.00 35.00
ATOM	1707 CB TYR A 216		31.763	21.683 114.753	1.00 36.19
ATCM	1708 CG TYR A 216		30.928	22.671 115.547	1.00 36.68
MOTA	1709 CD1 TYR A 216		29.961	23.462 114.925	1.00 35.83
ATOM	1710 CE1 TYR A 216		29.249	24.431 115.641	1.00 40.89
MOTA	1711 CD2 TYR A 216		31.163	22.869 116.910	1.00 41.50
ATOM	1712 CE2 TYR A 216		30.459	23.834 117.634	1.00 40.69
ATOM	1713 CZ TYR A 216		29.505	24.612 116.994	1.00 40.17
ATOM	1714 OH TYR A 216		28.816	25.566 117.708	1.00 38.09
ATOM	1715 C TYR A 216 1716 O TYR A 216		33.877		1.00 34.05
MOTA	1716 O TYR A 216		34.263	20.462 114.127	1.00 31.87

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ATO	OM 1785 CA LEU A 226	57 24.26
ATC	OM 1786 CB LEU A 226	1.00 25.35
ATC	OM 1787 CG LEU A 226	55 150 23,23
ATO	OM 1788 CD1 LEU A 226	2.00.20.20
ATC	M 1789 CD2 LEU A 226	2. 22.
ATC	M 1790 C LEU A 226	50 777
ATC	M 1791 O LEU A 226	2.00 24.75
ATO	M 1792 N ASN A 227	58.902 20.450 107.299 1.00 26.15
ATO		38.664 22.674 107.548 1.00 23.94
ATO		59.702 22.733 108.578 1.00 24.89
ATO	1.011 11 22 /	60.751 23.823 108.269 1.00 27.43
ATO		25.231 108.334 1.00 31.62
ATO	M 1797 ND2 ASN A 227	39.398 25.632 109.336 1.00 31.17
ATO		60.395 25.998 107.267 1.00 28.06
ATO		39.076 22.957 109.960 1.00 23.16
ATO		37.8/3 23.206 110.065 1.00 18 45
ATO		22.862 111.018 1.00 21 29
ATO	M 1802 CB ASP A 228	59.357 23.032 112.375 1.00 25.80
ATO	220	00.464 22.893 113.426 1.00 24.02
ATO		61.110 21.520 113.422 1.00 26.48
ATON		60.410 20.530 113.135 1.00 29 55
ATON		62.311 21.425 113.744 1.00 29 88
ATOM	220	58.628 24.341 112.620 1.00 27 83
ATOM		37.389 24.360 113.284 1.00 25.68
ATOM	***************************************	59.167 25.437 112.098 1.00 25.78
ATOM		58.537 26.739 112.297 1 00 27 75
ATOM		59.453 27.850 111.770 1.00 32.77
ATOM		28.020 112.621 1.00 30.35
ATOM	11 223	00.035 28.433 113.782 1.00 33.12
ATOM	11 127	61.856 27.691 112.053 1.00 28.11
ATOM		57.168 26.817 111.645 1.00 29.47
ATOM		56.230 27.387 112.202 1.00 26 75
ATOM		37.041 26:228 110.463 1.00 30.80
ATOM		55.761 26.244 109.773 1.00 30 77
ATOM		55.929 25.716 108.341 1.00 29.11
ATOM	220	56.897 26.531 107.507. 1.00 35.94
ATOM	1821 OE1 GLU A 230	57.119 25.946 106.125 1.00 37.77
MOTA	1822 OE2 GLU A 230	57.465 24.748 106.039 1.00 38.98
ATOM	1823 C GLU A 230	56.957 26.684 105.129 1.00 31.32
ATOM	1824 O GLU A 230	54.723 25.407 110.527 1.00 30.13
ATOM	1825 N PHE A 231	53.563 25.799 110.631 1.00 28.35
ATOM	1826 CA PHE A 231	55.141 24.262 111.060 1.00 32.49
ATOM	1827 CB PHE A 231	54.223 23.386 111.790 1.00 28.54
ATOM	1828 CG PHE A 231	54.913 22.075 112.191 1.00 31.22 53.974 21.050 112 781 1.00 38.41
ATOM	1829 CD1 PHE A 231	11.00 48.41
ATOM	1830 CD2 PHE A 231	1.00 29.00
ATOM	1831 CE1 PHE A 231	1.00 20.30
ATOM	1832 CE2 PHE A 231	En 166 10 23.73
ATOM	1833 CZ PHE A 231	1.00 31.40
ATOM	1834 C PHE A 231	2.00 50.51
ATOM	1835 O PHE A 231	20.00
MOTA	1836 N LEU A 232	1.00 23.39
MOTA	1837 CA LEU A 232	2.00 20.75
ATOM	1838 CB LEU A 232	55 100 27.25
ATOM	1839 CG LEU A 232	15. 15.
ATOM	1840 CD1 LEU A 232	56.176 24.372 116.420 1.00 28.11
ATOM	1841 CD2 LEU A 232	57.440 24.783 117.162 1.00 27.32
ATOM	1842 C LEU A 232	55.268 23.540 117.328 1.00 27.87
ATOM	1843 O LEU A 232	53.371 26.542 114.800 1.00 26.98
ATOM	1844 N PHE A 233	52.449 26.866 115.544 1.00 23.34
ATOM	1845 CA PHE A 233	53.694 27.232 113.708 1.00 24.99
ATCM ·	1646 CB PHE A 233	52.950 28.426 113.312 1.00 28.13
ATOM	1847 CG PHE A 233	53.542 29.029 112.029 1.00 30.77
ATOM	1948 CD1 PHE A 233	52.719 30.151 111.448 1.00 29.65 52.803 31.441 111.962 1.00 32.80
		52.803 31.441 111.962 1.00 32.80

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ATOM	1849	CI	2. PHE A 2	33	51.825	29.903	3 110.412	1.00 31.59
ATOM	1850) CE	E1 PHE A 2	33	52.008		3 111.452	
MOTA	1851	L CE	22 PHE A 2	33	51.022	30.924	109.895	
ATOM	1852	CZ	PHE A 2	33	51.114	32.208	3 110.415	
ATOM	1853	C	PHE A 2	33	51.510	27.999	113.031	1.00 31.62
MOTA	1854	0	PHE A 2	33	50.553	28.603	3 113.532	1.00 25.88
MOTA	1855	N	ALA A 2	14	51.370		112.215	
MOTA	1856	CA	ALA A 2	14	50.056		5 111.853	
ATOM	1857	CE	ALA A 2	4	50.195		110.864	1.00 20.08
MOTA	1858	C	ALA A 23		49.304		113.089	1.00 25.17
ATOM	1859	0	ALA A 23	4	48.114		113.234	1.00 25.21
ATOM	1860	N	LEU A 23		50.002	•	113.987	1.00 28.18
ATOM	1861	CA	LEU A 23	5	49.367		115.195	1.00 33.70
MOTA	1862	CB	LEU A 23	5	50.356		116.026	1.00 32.70
ATOM	1863	CG	LEU A 23	5	49.772	22.788	116.820	1.00 36.89
ATOM	1864	CD	1 LEU A 23	5	50.634	22.545	118.052	1.00 31.37
ATOM	1865	CD			48.344	23.072	117.231	1.00 31.47
ATOM	1866	С	LEU A 23	5 .	48.841	25.925	116.062	1.00 33.38
MOTA	1867	О	LEU A 23	5	47.673	25.926	116.455	1.00 28.13
ATOM	1868	N	GLU A 23		49.710	26.888	116.362	1.00 34.02
ATOM	18,69	CA	GLU A 23	6	49.336	28.026	117.199	1.00 37.30
MOTA	1870		GLU A 23		50.528	28.972	117.400	1.00 41.51
MOTA	1871	CG	GLU A 23		51.675		118.188	1.00 49.54
ATOM	1872	CD	GLU A 23				118.451	1.00 55.02
ATOM	1873	OE.			53.781		119.140	1.00 56.19
ATOM	1874	OE:			52.735		117.968	1.00 54.84
ATOM	1875	C	GLU A 23		48.163		116.638	1.00 33.98
ATOM	1876	O N	GLU A 23 LYS A 23		47.211		117.362	1.00 37.01
ATOM	1877 1878	CA	LYS A 23		48.223 47.140		115.354	1.00 33.94
ATOM ATOM	1879	CB	LYS A 23		47.505		114.726 113.281	1.00 33.10 1.00 36.08
ATOM	1880	CG	LYS A 23		48.695		113.261	1.00 38.08
ATOM	1881	CD	LYS A 23		48.395		113.103	1.00 33.02
ATOM	1882	CE	LYS A 23		49.569		113.762	1.00 45.24
ATOM	1883	NZ	LYS A 23		49.285		114.500	1.00 43.49
ATOM	1884	C	LYS A 23		45.820		114.751	1.00 31.40
ATOM	1885	0	LYS A 23	7	44.793		115.131	1.00 31.67
ATOM	1886	N	SER A 23	3	45.841	27.861	114.354	1.00 28.72
ATOM	1887	CA	SER A 23	3	44.610	27.080	114.335	1.00 31.74
ATOM .	1888	CB	SER A 23		44.834	25.720	113.660	1.00 28.90
ATOM	1889	OG	SER A 23		45.760		114.372	1.00 25.18
ATOM	1890	С	SER A 23		44.041		115.740	1.00 33.23
MOTA	1891	0	SER A 23		42.823		115.916	1.00 34.79
ATOM	1892	N	LEU A 23		44.907		116.741	1.00 35.27
ATOM	1893	CA	LEU A 239		44.413		118.108	1.00 37.57
MOTA	1894	CB	LEU A 239		45.554		119.090	1.00 38.58
ATOM	1895 1896	CG	LEU A 239		46.176		119.038	1.00 39.74
ATOM	1897		LEU A 239		47.276 45.109		120.075 119.301	1.00 35.82 1.00 34.93
ATOM	1898	CDZ	LEU A 239		43.109		118.521	1.00 34.93
ATOM	1899		LEU A 239		42.628	27.032	110.321	1.00 35.50
ATOM	1900	N	GLU A 240		44.202		118.131	1.00 33.30
ATOM ATOM	1901	CA	GLU A 240		43.561		118.450	1.00 40.15
ATOM	. 1902	CB	GLU A 240		44.366		117.883	1.00 40.42
ATOM	1903	CG	GLU A 240		45.661		118.602	1.00 43.91
ATOM	1904	CD	GLU A 240		46.407		117.942	1.00 49.31
ATOM	1905		GLU A 240		45.772	33,925	117.665	1.00 49.00
ATOM	1906	OE2	GLU A 240		47.624		117.705	1.00 54.05
ATOM	1907	c	GLU A 240		42.165		117.849	1.00 39.58
ATOM	1908	0	GLU A 240		41.224		118.455	1.00 40.99
ATCM	1909	N	ILE A 241		42.039	29.764	116.645	1.00 35.70
ATOM	1910	CA	ILE A 241		40.754	29.726	115.964	1.00 38.23
ATOM	1911	CB	ILE A 241		40.904	29.150	114.546	1.00 37.55
ATOM	1912	CG2	ILE A 241		39.535	29.005	113.895	1.00 37.30
ATOM	1913	CG1	ILE A 241		41.832	30.048	113.724	1.00 38.36
ATOM	1914	CD1	ILE A 241		42.106	29.541	112.320	1.00 36.15

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ATOM	1915 C ILE A 241	39	.751	28 85	31 116.73	7 1 0	0 37 3.
ATOM			.591		4 116.88		0 37.31
ATOM			.203				0 37.91
ATOM			.336	_			0 37.07
ATOM							0 38.35
			.025) 1.0	0 37.58
ATOM			.120			3 1.0	0 31.91
ATOM		40	.364	24.80	3 116.928	3 1.00	0 39.21
ATOM		38	.930	27.45	1 119.309	1.00	0 40.40
MOTA		37	.759	27.42			38.19
ATOM	1924 N LYS A 243		. 905		8 120.014		0 45.47
ATOM	1925 CA LYS A 243		661	28.65		1 00	0 40 74
ATOM	1926 CB LYS A 243		.945	29.31		. 1.00	48.74
ATOM	1927 CG LYS A 243		853			1.00	51.01
ATOM					6 122.614		56.09
			250	28.14			57.39
MOTA			054		6 124.783		59.53
ATOM	1930 NZ LYS A 243		448		3 126.127	1.00	57.91
ATOM	1931 C LYS A 243		559	29.70		1.00	51.67
ATOM	1932 O LYS A 243	· 37.	815	29.87	1 122.226	1.00	52.84
MOTA	1933 N GLU A 244	38.	451		0 120.140		53.77
ATOM	1934 CA GLU A 244	37.	460		1 120.004		54.74
ATOM	1935 CB GLU A 244		954		7 118.986		55.15
MOTA	1936 CG GLU A 244	37.	068		7 118.865	1.00	60.63
ATOM	1937 CD GLU A 244		602		117.868	1.00	60.63
ATOM	1938 OE1 GLU A 244		746			4.00	65.87
ATOM	1939 OE2 GLU A 244				118.053		70.36
ATOM	1940 C GLU A 244		879		116.900	1.00	67.09
ATOM		36.			119.626		52.65
		35.		31.838	3 119.606	1.00	55.59
ATOM	1942 N VAL A 245	35.			119.332	1.00	50.57
ATOM	1943 CA VAL A 245	34.			118.947	1.00	45.75
ATOM	1944 CB VAL A 245	34.	475	29.081	117.409	1.00	46.91
MOTA	1945 CG1 VAL A 245	33.	085	28.634	116.986		52.62
ATOM	1946 CG2 VAL A 245	34.	825	30.389	116.716	1.00	48.34
MOTA	1947 C VAL A 245	34.	130		119.642	1.00	42.67
MOTA	1948 O VAL A 245	33.0			119.445		43.61
ATOM	1949 N PHE A 246	35.0			120.477	1 00	40.87
MOTA	1950 CA PHE A 246	34.6			121.139	1.00	37.47
ATOM	1951 CB PHE A 246	35.3			120.257	1.00	37.47
ATOM	1952 CG PHE A 246	34.4		23.685			
ATOM	1953 CD1 PHE A 246	33.3					33.22
ATOM	1954 CD2 PHE A 246			23.467			33.93
MOTA	1955 CE1 PHE A 246	35.1		22.674			32.13
ATOM	1956 CE2 PHE A 246	32.4			120.621		37.75
		34.5		21.459			35.92
MOTA		33.2		21.252	121.251		36.30
ATOM	1958 C PHE A 246	35.3		26.065	122.509	1.00	38.93
ATOM	1959 O PHE A 246	36.5		26.158	122.630	1.00	40.66
MOTA	1960 N GLU A 247	34.5	00	25.870	123.537		38.59
MOTA	1961 CA GLU A 247	34.9	70	. 5.733	124.918		44.60
ATOM	1962 CB GLU A 247	34.1	46	: 5.615	125.865		47.07
ATOM	1963 CG GLU A. 247	33.1	61	27.569	125.185		56.16
ATOM	1964 CD GLU A 247	31.9	44	26.865	124.577		62.03
ATOM	1965 OE1 GLU A 247	32.0			123.607		
ATOM	1966 OE2 GLU A 247	30.8			125.083		61.85
ATOM	1967 C GLU A 247						64.59
	1968 0 GLU A 247	34.7			125.285		39.40
ATOM		33.7			125.794		39.91
MOTA	1969 N PRO A 248	35.7			125.041	1.00	
ATOM	1970 CD PRO A 248	37.1			124.483	1.00	3325
	1971 CA PRO A 248	35.7			125.316	1.00	35.84
MOTA	1972 CB PRO A 248	37.0		21.531	124.648	1.00	
ATOM	1973 CG PRO A 248	37.91	70		124.982	1.00	
ATOM	1974 C PRO A 248	35.73		21.611	126.779	1.00	
ATOM	1975 0 PRO A 248	36.44			127.597	1.00	
ATOM	1976 N GLU A 249	34.91			127.096	1.00	20 20
ATOM	1977 CA GLU A 249	34.84			128.459	1.00	
ATOM ·	1978 CB GLU A 249	33.52			128.693	1 00	33.IX
atom	1979 CG GLU A 249	32.28				1.00	
	1980 CD GLU A 249	31.02			128.564	1.00	
atcm	CD GDO N 247	21.02		LJ.388	128.668	1.00 4	90.52

MOTA	1981	OE	1 GLU A	249	30.817	18.509	127.804	1.00 40.27
ATOM	1982				30.252		129.620	1.00 38.57
	1983		_				128.623	1.00 32.30
MOTA			GLU A		35.995			
MOTA	1984	0	GLU A	249	36.472	18.884	129.728	1.00 28.51
MOTA	1985	N	VAL A	250	36.434	18.547	127.502	1.00 33.74
ATOM.	1986	CA	VAL A	250	37.516	17 563	127.494	1.00 29.31
	1987		VAL A					
ATOM					36.988		127.926	1.00 29.85
MOTA	1988	CG	1 VAL A		35.908	15.711	126.958	1.00 24.36
ATOM	1989	CG:	2 VAL A	250	38.121	15.166	127.978	1.00 25.60
ATOM	1990	C.	_VAL A	250	38.066	17.453	126.076	1.00 29.30
	1991		VAL A		37.358		125.114	1.00 24.46
ATOM								
ATOM	1992	N	TYR A		39.323		125.930	1.00 27.96
MOTA	1993	CA	TYR A	251 .	39.865	16.913		1.00 30.06
ATOM	1994	CB	TYR A	251	40.585	18.206	124.165	1.00 25.89
ATOM	1995	CG	TYR A	251	41.998		124.692	1.00 29.90
	1996	CD:			43.087		124.029	1.00 26.02
MOTA								
ATOM	1997	CE:			44.390	17.953		1.00 29.20
MOTA	1998	CD	2 TYR A	251	42.249	19.107	125.849	1.00 31.96
MOTA	1999	CE	2 TYR A	251	43.551	19.271	126.338	1.00 31.54
ATOM	2000	CZ	TYR A	251	44.614	18.694	125.664	1.00 31.46
	2001	ОН	TYR A		45.894		126.152	1.00 29.69
MOTA								1.00 27.56
MOTA	2002	С	TYR A		40.801		124.451	
MOTA	2003	0	TYR A		41.382		125.436	1.00 28.23
ATOM	2004	N	LEU A	252	40.908	15.222	123.227	1.00 23.52
ATOM	2005	CA	LEU A	252	41.806	14.117	122.919	1.00 26.53
MOTA	2006	CB	LEU A		41.057		122.293	1.00 25.74
	2007	CG	LEU A		40.266		123.221	1.00 28.49
ATOM								
MOTA	2008	CD1			39.122	12.753		1.00 27.67
MOTA	2009	CD2			39.72 7		122.414	1.00 32.00
MOTA	2010	С	LEU A	252	42.842	14.638	121.932	1.00 27.53
ATOM	2011	0	LEU A	252	42.528	15.444	121.055	1.00 24.42
ATOM	2012	N	LEU A		44.075	14.176	122.078	1.00 24.60
ATOM	2013	CA	LEU A		45.157		121.204	1.00 25.04
	2014	CB	LEU A		46.176		122.017	1.00 22.48
ATOM							121.323	1.00 21.05
MOTA	2015	CG	LEU A		47.456			
ATOM	2016		LEU A		47.105	16.833	120.175	1.00 23.05
ATOM	2017	CD2			48.348		122.360	1.00 16.40
ATOM	2018	С	LEU A	253	45.822	13.374	120.580	1.00 23.55
ATOM	2019	0	LEU A	253	46.329	12.516	121.303	1.00 22.11
MOTA	2020	N	GLN A	254	45.811	13.287	119.248	1.00 22.33
	2021	CA	GLN A		46.417		118.552	1.00 19.84
MOTA	2022		GLN A		45.542		117.348	1.00 23.09
MOTA		CB						
ATOM	2023	CG	GLN A		46.075		115.963	1.00 35.49
ATOM	2024	CD	GLN A		47.073		115.453	1.00 31.26
ATOM '	2025	OE1	GLN A	254	46.712	9.937	114.961	1.00 33.69
: TOM	2026	NE2	GLN A	254	48.338	11.349	115.574	1.00 31.02
	2027	c	GLN A		47.831		118.153	1.00 22.46
. TOM -							117.478	1.00 17.56
FOM	2028	0	GLN A		48.034	13.599	117.470	
ATOM	2029	N	LEU A		48.804	11.781	118.590	1.00 17.64
ATOM	2030	CA	LEU A		50.213	12.079	118.383	1.00 17.04
MOTA	2031	CB	LEU A	255	50.894	12.136	119.750	1.00 14.75
	2032	CG	LEU A	255	50.277		120.670	1.00 25.02
ATOM				255	50.732	12 996	122.107	1.00 21.99
ATOM	2033	CD1				12.330	122.107	
ATOM	2034		LEU A		50.636	14.5/8	120.149	1.00 18.30
ATOM	2035	С	LEU A		51.023	11.169	117.476	1.00 21.34
ATOM	2036	0	LEU A	255	52.089	10.705	117.875	1.00 18.73
ATOM	2037	N	GLY A		50.543	10.928	116.259	1.00 22.75
	2038	CA	GLY A		51.291	10 093	115.330	1.00 24.09
MOTA			GLY A				115.126	1.00 24.27
MOTA	2039	C			52.660			
MOTA	2040	0	GLY A		52.805	11.945	115.134	1.00 19.15
ATOM	2041	Ŋ	THR A	257	53.680	9.903	114.948	1.00 24.14
ATOM	2042	CA	THR A		55.014		114.765	1.00 21.32
ATOM	2043	CB	THR A	25 7	56.048	9.582	115.511	1.00 17.24
	2044	0G1	THR A	257	56.009		115.004	1.00 17.48
ATOM	_	CG2	THR A	257	55.728	9.532	117.016	1.00 15.73
ATOM	2045		THR A	257		3.332	113.290	1.00 22.98
ATOM	2046	C	100 2		55.403	10.54/	220.270	2.00 22.90

	•	_		
ATO	11 23/	56.517	7 10.941 112.974 1.00 20	
ATO	230	54.493	10 100 20	.39
ATO		54.863	10 000 000	.56
ATO	11 230	53.849	0 405 5	.06
ATO		52.415		.08
ATO	250	55.222	11.596 110.364 1.00 27	
ATON		55.756	11.661 109.254 1.00 25	61
ATON		52.173	11.070 110.742 1.00 29	.86
ATON		51.513	9.168 109.869 1.00 33	.25
ATOM ATOM	-110 11 200	54.884	12.710 111.045 1.00 31	.06
ATOM		54.019	12.928 112.220 1.00 28	.59
· ATOM		55.268	14.006 110.469 1.00 30	.95
ATOM		54.447	14.993 111.296 1.00 35	.06
ATOM	2061 C PRO A 259	54.418	14.320 112.636 1.00 35.	.26
ATOM		56.790	14.268 110.583 1.00 29	.06
ATOM	200	57.300 57.508	15.251 110.044 1.00 29	.70
ATOM	200	58.960	13.389 111.280 1.00 22.	. 68
ATOM		59.461	13.545 111.455 1.00 28. 12.576 112.533 1.00 22	41
ATOM	2066 CG LEU A 260	58.970		47
ATOM	2067 CD1 LEU A 260	59.352		14
ATOM	2068 CD2 LEU A 260	59.592	11.599 114.826 1.00 22. 14.079 114.532 1.00 20.	83
MOTA	2069 C LEU A 260	59.770	13.344 110.160 1.00 27.	48
MOTA	2070 O LEU A 260	59.407	12.535 109.299 1.00 27.	72
ATOM	2071 N LEU A 261	60.874	14.081 110.040 1 00 26	76
ATOM	2072 CA LEU A 261	61.742	14.010 108.865 1.00 26	56
MOTA MOTA	2073 CB LEU A 261 2074 CG LEU A 261	63.067	14.737 109.137 1.00 23	06
ATOM		64.131	14.615 108.025 1 00 29	52
ATOM	2075 CD1 LEU A 261 2076 CD2 LEU A 261	63.642	15.325 106.770 1.00 22.	68
ATOM	2077 C LEU A 261	65.460	15.219 108.475 1.00 26.	71
ATOM	2078 O LEU A 261	62.063 61.880	12.577 108.443 1.00 28.	23
ATOM	2079 N GLU A 262	62.539	12.198 107.289 1.00 26.	52
ATOM	2080 CA GLU A 262	62.938	11.787 109.397 1.00 28.10.416 109.135 1.00 31.	70
ATOM	2081 CB GLU A 262	63.685		76
ATOM	2082 CG GLU A 262	64.890	9.855 110.351 1.00 29.3 10.683 110.803 1.00 31.3	12
ATOM	2083 CD GLU A 262	64.521	11.847 111.708 1.00 28.0	
MOTA	2084 OE1 GLU A 262	63.324	12.195 111.789 1.00 28.7	
ATOM	2085 OE2 GLU A 262	65.433	12.424 112.340 1.00 26.0	
ATOM	2086 C GLU A 262	61.847	9.429 108.721 1.00 28.6	
MOTA MOTA	2087 O GLU A 262 2088 N ASP A 263	62.158	8.305 108.350 1.00 29.7	12
ATOM	2088 N ASP A 263 2089 CA ASP A 263	60.582	9.825 108.785 1.00 28.0	7
ATOM	2090 CB ASP A 263	59.513	8.902 108.412 1.00 26.8	
ATOM	2091 CG ASP A 263	58.305 57.261	9.099 109.333 1.00 25.2	6
ATOM	2092 OD1 ASP A 263	56.638	7.998 109.185 1.00 33.1	.4
ATOM	2093 OD2 ASP A 263	57.042	7.636 110.209 1.00 29.9 7.509 108.051 1.00 26.5	
ATOM	2094 C ASP A 263	59.150		5
ATOM	2095 O ASP A 263			4
ATOM	2096 N TYR A 264	59.303	10.247 106.594 1.00 24.7 8.111 106.130 1.00 27.5	0
ATOM	2097 CA TYR A 264	59.031	8.219 104.696 1.00 33.8	Ţ
ATOM	2098 CB TYR A 264	59.576	7.008 103.935 .1.00 40.4	9 1
ATOM	2099 CG TYR A 264	61.059	6.771 104.092 1.00 50.6	a A
ATOM	2100 CD1 TYR A 264	61.565	6.087 105.199 1.00 54.6	7
MOTA	2101 CE1 TYR A 264	62.933	5.876 105.351 1.00 55.28	B
ATOM	2102 CD2 TYR A 264 2103 CE2 TYR A 264	61.960	7.242 103.140 1.00 53.75	
MOTA MOTA		63.329	7.038 103.282 1.00 56.61	Ŀ
ATOM		63.809	6.354 104.388 1.00 56.22	2
ATOM	2105 OH TYR A 264 2106 C TYR A 264	65.161 57.581	6.147 104.524 1.00 55.90)
ATOM	2107 O TYR A 264	57.381 57.311	8.394 104.294 1.00 31.33	
ATOM	2108 N LEU A 265	56.641	8.825 103.178 1.00 27.15	
MOTA	2109 CA LEU A 265	55.244	8.059 105.172 1.00 26.88	
MOTA	2110 CB LEU A 265	54.360	8.209 104.792 1.00 24.39 7.189 105.527 1.00 26.55	
MOTA	2111 CG LEU A 265	54.663		
MOTA	2112 CD1 LEU A 265			
_		J J- 3	4.836 105.503 1.00 21.17	

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ATOM	2113	CD2 LEU A 265	54.931	5.620 103.682	1.00 33.35
ATOM	2114	C LEU A 265	54.669		1.00 20.81
ATOM	2115		53.457	9.796 104.979	1.00 21.30
MOTA	2116		55.540	10.622 104.959	1.00 23.23
MOTA	2117		55.084		1.00 26.30
ATOM	2118		54.856		1.00 25.16
ATOM	2119		56.074		1.00 22.92
MOTA	2120		56.147		1.00 30.17
MOTA	2121		57.334		1.00 31.65
ATOM	2122 2123		55.731		1.00 31.56
ATOM ATOM	2123		56.696		1.00 27.65
ATOM	2125		56.140 55.815		1.00 30.54
ATOM	2126		57.039		1.00 34.13 1.00 29.09
ATOM	2127		56.745	12.376 99.524	1.00 23.03
ATOM	2128		57.956	11.541 99.272	1.00 37.01
MOTA	2129	C LYS A 267	57.050	16.004 104.107	1.00 30.85
MOTA	2130	O LYS A 267	57.624	17.017 103.707	1.00 27.86
MOTA	2131	N PHE A 268	56.688	15.826 105.377	1.00 24.19
MOTA	2132	CA PHE A 268	57.009	16.808 106.412	1.00 25.34
ATOM	2133	CB PHE A 268	56.014	16.730 107.579	1.00 24.54
ATOM	2134	CG PHE A 268	54.636	17.256 107.257	1.00 21.68
MOTA	2135	CD1 PHE A 268	53.631	17.221 108.216	1.00 28.65
MOTA	2136	CD2 PHE A 268 CE1 PHE A 268	54.346	17.806 106.011	1.00 25.14
ATOM ATOM	2137 2138	CE1 PHE A 268 CE2 PHE A 268	52.357	17.728 107.944	1.00 25.34
ATOM	2139	CZ PHE A 268	53.077 52.082	18.315 105.730 18.275 106.702	1.00 23.79
ATOM	2140	C PHE A 268	58.410	18.275 106.702 16.470 106.908	1.00 28.13 1.00 25.66
ATOM	2141	O PHE A 2.68	58.778	15.299 106.994	1.00 28.44
ATOM	2142	N ASN A 269	59.194	17.490 107.230	1.00 25.81
MOTA	2143	CA ASN A 269	60.555	17.270 107.709	1.00 30.60
ATOM	2144	CB ASN A 269	61.566	17.938 106.767	1.00 31.97
MOTA	2145	CG ASN A 269 ·		17.513 105.317	1.00 35.83
MOTA	2146	OD1 ASN A 269	61.235	16.332 105.020	1.00 33.93
ATOM	2147	ND2 ASN A 269	61.446	18.477 104.405	1.00 33.95
ATOM	2148 2149	C ASN A 269 O ASN A 269	60.723	17.843 109.110	1.00 31.80
atom atom	2150	N LEU A 270	61.609 59.888	18.665 109.348 17.397 110.043	1.00 28.13 1.00 29.70
ATOM	2151	CA LEU A 270	59.954	17.918 111.406	1.00 25.70
ATOM	2152	CB LEU A 270	58.575	17.833 112.074	1.00 26.60
ATOM	2153	CG LEU A 270	57.392	18.425 111.297	1.00 29.62
ATOM	2154	CD1 LEU A 270	56.177	18.494 112.222	1.00 28.54
MOTA	2155	CD2 LEU A 270	57.740	19.825 110.790	1.00 29.40
ATOM	2156	C LEU A 270	60.979	17.242 112.301	1.00 26.83
ATOM	2157	O LEU A 270	61.490	16.158 111.990	1.00 19.60
MOTA	2158	N SER A 271	61.275	17.896 113.420	1.00 21.66
ATOM ATOM	2159 2160	CA SER A 271 CB SER A 271	62.220 63.189	17.365 114.393	1.00 27.08
ATOM	2161	OG SER A 271	62.499	18.460 114.846 19.433 115.626	1.00 24.64
ATOM	2162	C SER A 271	61.454	16.868 115.618	1.00 18.60
ATOM	2163	O SER A 271	60.272	17.150 115.772	1.00 22.56
ATOM	2164	N ASN A 272	62.157	16.129 116.470	1.00 28.35
MOTA	2165	CA ASN A 272	61.649	15.593 117.739	1.00 31.03
ATOM	2166	CB ASN A 272	62.774	14.880 118.498	1.00 28.80
ATOM	2167	CG ASN A 272	62.854	13.428 118.180	1.00 36.65
ATOM	2168	OD1 ASN A 272	63.712	12.709 118.705	1.00 29.89
ATOM	2169	ND2 ASN A 272	61.953	12.969 117.319	1.00 40.80
ATOM	2170	C ASN A 272	61.167	16.695 118.661	1.00 31.83
ATOM	2171	O ASN A.272 N VAL A 273	60.090	16.618 119.261	1.00 27.50
ATOM	2172 2173	CA VAL A 273	62.032 61.802	17.693 118.804	1.00 31.49
ATOM ATOM	2174	CB VAL A 273	63.069	18.837 119.667 19.709 119.725	1.00 31.38
ATOM	2175	CG1 VAL A 273	62.804	20.988 120.500	1.00 48-07
ATOM	2176	CG2 VAL A 273	64.198	18.914 120.381	1.00 42.81
ATCM	2177	C VAL A 273	60.608	19.665 119.234	1.00 30.13
ATOM	2178	O VAL A 273	59.872 .	20.174 120.072	1.00 31.44

		. •
ATO	M 2179 N ALA A 274	60 405 10 000
ATO		60.405 19.800 117.929 1.00 24.15
ATO		39.238 20.558 117.455 1.00 26 27
ATO		59.341 20.780 115.965 1.00 21.85
	x 2/4	
ATO		
ATO	M 2184 N PHE A 275	50 450 45.76
ATO	M 2185 CA PHE A 275	
ATO		37.015 17.538 117.974 1.00 25 gg
ATO	A Z/J	37.449 16.092 117.710 1.00 25 21
		56.340 15.088 117.870 1.00 28.85
ATO		Fr 070
ATO	A Z/J	
ATON	M 2190 CE1 PHE A 275	20,33
OTA	1 2191 CE2 PHE A 275	EF 3.5
ATON	1 2192 CZ PHE A 275	33.343 13.231 119.059 1.00 30 83
ATOM		54.282 13.214 118.160 1.00 34 19
		56.607 17.712 119 445 1 00 24 65
ATOM		EE 400 2.00 24.03
ATOM	, 220 A 2/0	52 50 40
ATOM	1 2196 CA LEU A 276	FR 7
ATOM	2197 CB LEU A 276	50 55-
ATOM		30.00/ 1/.692 122.534 1 00 26 11
ATOM		30.031 18.132 124.001 1 00 31 15
		57.609 17.351 124.761 1 00 20 22
MOTA	320 2 2/0	60 033 37 037 101
ATOM	2201 C LEU A 276	56 555
ATOM	2202 O LEU A 276	EE 000 10 11 12
ATOM	2203 N LYS A 277	
ATOM	220 72 277	37.333 20.219 121.425 1.00 30 99
ATOM	/1 2//	56.913 21.593 121 603 1 00 27 04
		57.742 22.516 120 704 1 00 30 30
ATOM	2206 CG LYS A 277	TE 0.00 JU.JO
MOTA	2207 CD LYS A 277	56 65 50 50 40
MOTA	2208 CE LYS A 277	FC 1.00 42./3
ATOM	2209 NZ LYS A 277	1.00 4J./U
ATOM	2210 C LYS A 277	57.528 26.004 123.390 1.00 44.64
ATOM		33.432 21.683 121.242 1.00 30 26
ATOM		54.640 22.284 121.972 1.00 27 55
		55.057 21.078 120.115 1.00 30.15
ATOM	2213 CA ALA A 278	
MOTA	2214 CB ALA A 278	E3 406 00 30.31
ATOM	2215 C ALA A 278	E2 700 00 TO
ATOM	2216 O ALA A 278	
MOTA	2217 N PHE A 279	51.735 21.067 121.108 1.00 30.36
ATOM	2218 CA PHE A 279	33.445 19.422 121.360 1.00 27 g5
ATOM	2219 CB PHE A 279	52.540 18.759 122,448 1.00 29 62
	11 2/2	53.343 17.534 122.886 1 00 26 02
ATOM	2220 CG PHE A 279	52.786 16.823 124.078 1.00 29.11
ATOM	2221 CD1 PHE A 279	C. 27.11
ATOM	2222 CD2 PHE A 279	1.00 20.00
MOTA	2223 CE1 PHE A 279	F4 0-1
ATOM	2224 CE2 PHE A 279	51.054 15.500 125.121 1.00 37.90
ATOM	THE R 2/3	. 53.011 16.114 126.386 1.00 38.01
ATOM	2/3	51.783 15.469 126 313 1 00 36 1
		52.362 19.730 123.621 1.00 30.67
ATOM	2227 O PHE A 279	
ATOM	2228 N ASN A 280	1.00 20.20
ATOM	2229 CA ASN A 280	
MOTA	2230 CB ASN A 280	53.339 21.373 125.104 1.00 30.54
	1.01. 11 200	34·/24 21.819 125.583 1.00 26 79
		33.508 20.695 126.227 1.0h 33 68
ATOM	2232 OD1 ASN A 280	54.958 19.886 126.971 1.00 34.47
ATOM	2233 ND2 ASN A 280	
ATOM	2234 C ASN A 280	53 400
MOTA	2235 O ASN A 280	C1 000
MOTA	2236 N ILE A 281	52 400
MOTA		32.429 22.960 123.509 1.00 27 32
		51.620 24.107 123.128 1.00 31.07
MOTA	2238 CB ILE A 281	51.878 24.517 121.666 1 00 35 00
ATOM	2239 CG2 ILE A 281	
ATOM .	2240 CG1 ILE A 281	E2 2E2 121.1/4 1.00 34.33
ATCM	2241 CD1 ILE A 281	E2 E20 == 1-00 JJ.JJ
	2242 C ILE A 281	50 144 5000 200 270 1.00 34.00
ATOM	2243 O ILE A 281	30.141 23.798 123.329 1.00 31.22
		49.391 24.621 123.854 1.00 30.15
TOM	2244 N VAL A 282	49.723 22.606 122.923 1.00 30.91
		1-00 JU.JI

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Figure 18-35

ATOM	2245	CA	VAL A 282		48.332	22.214	123.081	1.00	30.76
MOTA	2246	CB	YAL A 282		48.075	20.797	122.523		35.10
ATOM	2247		1 VAL A 282		46.641		122.841		28.72
ATOM	2248		2 VAL A 282		48.313		121.018		28.66
ATOM	2249		VAL A 282		47.952		124.558		31.39
ATOM	2250	0	VAL A 282		46.884	22.715	124.917	1.00	32.70
MOTA	2251	N	ARG A 283		48.837	21.720	125.406	1.00	29.86
ATOM	2252		ARG A 283		48.587	21.675	126.840		34.82
ATOM	2253	CB	ARG .A 283		49.629		127.519		31.44
_					49.551		127.061		
ATOM	2254	CG.							29.49
MOTA	2255	CD	ARG A 283		50.729		127.554		30.67
ATOM	2256	NE	ARG.A 283		50.730		128.990		30.78
ATOM	2257	CZ	ARG A 283	•	51.826	18.351	129.742	1.00	35.27
ATOM	2258	NH:	1 ARG A 283		53.012	18.611	129.198	1.00	36.46
ATOM	2259	NH	2 ARG A 283		51.742	18.100	131.035		35.90
ATOM	2260	С	ARG A 283		48.561		127.473		36.06
									•
ATOM	2261	0	ARG A 283		47.830		128.439		35.04
ATOM	2262	N	GLU A 284		49.350		126.928		35.70
ATOM	. 2263	CA	GLU A 284		49.376	25.348	127.448	1.00	40.93
MOTA	2264	CB	GLU A 284		50.499	26.166	126.799	1.00	44.17
ATOM	2265	CG	GLU A 284		51.917	25.702	127.141	1.00	56.39
ATOM	2266	CD	GLU A 284		52.989	26.495	126.401		60.69
ATOM	2267	OE1			53.012		126.542		63.13
	2268	OE2			53.810	•	125.680		62.79
ATOM									
ATOM	2269	C	GLU A 284		48.039		127.148		39.24
ATOM	2270	0	GLU A 284		47.525		127.954		38.52
MOTA	2271	N	VAL A 285		47.472	25.704	125.986		33.75
ATOM	2272	CA	VAL A 285		46.205	26.294	125.592	1.00	35.82
MOTA	2273	CB	VAL A 285		46.039	26.291	124.062	1.00	34.14
ATOM	2274	CG1	VAL A 285		44.654	26.811	123.693	1.00	36.43
ATOM	2275	CG2		•	47.114		123.419		37.26
ATOM	2276	c	VAL A 285		44.964		126.192		38.96
					44.043				
ATOM	2277	0	VAL A 285				126.611		41.83
ATOM	2278	N	PHE A 286		44.931		126.236		37.57
MOTA	2279	CA	PHE A 286		43.760		126.753		35.05
ATOM	2280	СB	PHE A 286		43.159		125.657		32.53
ATOM	2281	CG	PHE A 286		42.544	23.490	124.529	1.00	30.15
ATOM	2282	CD1	PHE A 286		43.104	23.459	123.256	1.00	33.96
ATOM	2283	CD2	PHE A 286		41.398	24.245	124.736	1.00	30.30
ATOM	2284	CE1			42.527		122.202		32.96
ATOM	2285	CE2			40.813		123.693		31.67
	2286	CZ	PHE A 286		41.381	24.919			31.66
ATOM									
MOTA	2287	C	PHE A 286		43.922		128.015		35.18
MOTA	2288	0	PHE A 286		42.984		128.409		36.97
MOTA	2289	N	GLY A 287		45.086	22.840	128.656	1.00	30.87
ATOM	2290	CA	GLY 7 287		45.297	22.056	129.862	1.00	30.06
ATOM	2291	C-	GLY 287		45.525	20.590	129.527	1.00	34.44
MOTA	2292	0	GLY 2. 287		45.914	20.264	128.403	1.00	32.54
MOTA	2293	N	GLU A 288		45.288		130.500	1.00	28.42
ATOM	2294	CA	GLU A 288		45.464		130.310		32.23
	2295	CB	GLU A 288				131.663		37.02
ATOM			020 000						
ATOM	2296	CG	GLU A 288		46.910		132.411		45.36
ATOM	2297	CD	GLU A 288		48.140		131.622		45.65
MOTA	2298	OE1	GLU A 288		48.144		131.069		46.42
MOTA	2299	OE2	GLU A 288		49.106	18.245	131.571	1.00	50.78
ATOM	2300	С	GLU A 288		44.309	17.623	129.546		30.98
ATOM	2301	o	GLU A 288		43.144	17.957	129.762		28.67
ATOM	2302	N	GLY A 289		44.641		128.657		29.66
			GLY A 289		43.625		127.886		28.08
MOTA	2303	CA							
ATOM	2304	C	GLY A 289		43.922		127.861		30.45
ATCM	2305	0	GLY A 289		44.618	14.307	128.739		25.90
ATOM	330é	И	VAL A 290		43.384	13.807	126.868	1.00	26.21
ATOM	2307	CA	VAL A 290		43.612	12.373	126.718		27.31
ATOM	2308	CB	VAL A 290		42.288	11.626	126.412	1.00	26.53
ATOM	2309		VAL A 290		42.554	10.148	126.204	1.00	25.36
ATOM	2310	CG2	VAL A 290		41.308	11.822	127.565		24.38
NI ON					,		-		

ATO	M 2311 C VAL A 290	44.580	3 3 3 4 0 1 0 5
ATO			
ATO		44.30	1 12.743 124.461 1 00 26 9
		45.71	9 11.597 125.775 1 00 23 5.
ATO		46.729	11.478 124.732 1.00 23.7
ATO		48.092	
lota	4 2316 CG TYR A 291	48.040	2.00 10.4
ATO			1.00 21.6
ATON	2318 CE1 TYR A 291	48.326	1.00 23.8
ATOM	2310 CD1 TIK A 291	48.200	14.331 128.214 1.00 24 5
		47.634	14.291 125.503 1 00 18 1-
ATOR		47.504	15.476 126.220 1.00 26.36
ATOM	1 2321 CZ TYR A 291	47.786	
ATOM	1 2322 OH TYR A 291	47.631	1.00 27.00
ATOM	2323 C TYR A 291		1.00 28.92
ATOM		46.768	1.00 23.03
	251	46.837	9.082 124.707 1.00 20 66
ATOM		46.755	10.142 122.711 1.00 23.96
ATOM	2326 CA LEU A 292	46.767	
ATOM	2327 CB LEU A 292	45.482	
ATOM		44.162	
ATOM	2329 CD1 LEU A 292	44.102	9.063 121.814 1.00 23.78
	2325 CD1 LEU A 292 .	43.001	8.959 120.826 1.00 23 09
ATOM		44.008	8.050 122.930 1.00 16.01
ATOM		47.953	8.885 120.947 1.00 22.90
ATOM	2332 O LEU A 292	. 48.527	
ATOM		48.301	
ATOM	2334 CA GLY A 293		7.684 120.491 1.00 18.83
ATOM	2335 C GLY A 293	49.401	7.529 119.554 1.00 24.35
ATOM		. 48.908	7.846 118.154. 1.00 24 22
		48.025	8.684 117.991 1.00 20 46
ATOM	2337 N GLY A 294	49.459	7.177 117.148 1.00 24.63
ATOM	2338 CA GLY A 294	49.035	
ATOM	2339 C GLY A 294	50.024	
ATOM	2340 O GLY A 294	50.956	
ATOM	2341 N GLY A 295		6.150 115.136 1.00 24.10
ATOM	233	49.825	7.203 113.499 1.00 19.85
		50.721	6.724 112.458 1.00 23.33
ATOM	2343 C GLY A 295	52.185	7.010 112.740 1.00 19.01
ATOM	2344 O GLY A 295 .	52.541	8.094 113.196 1.00 19.39
ATOM	2345 N GLY A 296	53.035	
ATOM	2346 CA GLY A 296	54.468	
ATOM	2347 C GLY A 296		6.162 112.690 1.00 22.65
ATOM	2348 O GLY A 296	55.098	4.898 112.146 1.00 25.31
		54.778	3.798 112.609 1.00 25.86
ATOM	2349 N TYR A 297	56.005	5.034 111.185 1.00 22.83
ATOM	2350 CA TYR A 297	56.598	3.852 110.577 1.00 23.93
ATOM	2351 CB TYR A 297	56.137	
ATOM	2352 CG TYR A 297	54.660	
ATOM	2353 CD1 TYR A 297	54.203	
ATOM	2354 CE1 TYR A 297		5.402 109.017 1.00 22.28
		52.842	5.695 109.089 1.00 20.19
ATOM	2355 CD2 TYR A 297	53.713	3.062 109.116 1.00 25.28
ATOM	2356 CE2 TYR A 297	52.352	3.346 109.190 1.00 21.83
ATOM	2357 CZ TYR A 297	51.927	
ATOM	2358 OH TYR A 297	50.588	
ATOM	2359 C TYR A 297	58.104	• • • • • • • • • • • • • • • • • • • •
ATOM .	2360 O TYR A 297	20.104	3.694 110.674 1.00 23.78
		58.665	2.724 110.154 1.00 22.07
ATOM		58.765	4.641 111.326 1.00 23.71
ATOM	2362 CA HIS A 298	60.204	4.534 111.517 1.00 26.33
ATOM	2363 CB HIS A 298	60.913	F 000 000
ATOM	2364 CG HIS A 298	62.403	+0.,3
ATOM	2365 CD2 HIS A 298	63.273	=:00 33.00
ATOM	2366 ND1 HIS A 298		5.465 112.215 1.00 31.83
	2300 NDI NIS A 298	63.151	5.775 110.056 1.00 32.49
ATOM	2367 CE1 HIS A 298	64.419	.5.547 110.345 1.00 29.10
ATCM	2368 NE2 HIS A 298	64.520	5.354 111.648 1.00 38.70
ATOM	2369 C HIS A 298	60.371	
ATOM	2370 O HIS A 298	60.120	
ATOM	2371 N PRO A 299		5.020 113.865 1.0C 25.07
		60.829	2.963 113.297 1.00 29.37
ATOM	2372 CD PRO A 299	61.285	1.932 112.352 1.00 26.09
rtom	2373 CA PRO A 299	61.024	2.491 114.669 1.00 29.46
ATOM	2374 CB PRO A 299	61.675	1.118 114.465 1.00 29.03
ATOM	2375 CG PRO A 299	62.411	
ATOM	2376 C PRO A 299	61.849	
0.1	2-1-2 - 110 11 202	01.043	3.403 115.570 1.00 31.88

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ATOM	237	7 O PROA 299	61.480	3.622 116.724	1.00 32.45
ATOM	2378		62.959		
MOTA	2379		63.803		
ATOM	2380		65.163	5.058 115.207	
ATOM	2381				
			65.912	3.821 114.738	
ATOM	2382		65.517	2.539 115.120	
ATOM	2383		66.214	1.411 114.682	1.00 30.68
MOTA	2384		67.027	3.941 113.908	1.00 29.72
ATOM	2385	CE2 TYR A 300	67.730	2.829 113.466	
MOTA	2386	CZ TYR A 300	67.320	1.568 113.854	1.00 33.89
ATOM	2387		68.011	0.471 113.404	1.00 34.70
ATOM	2388		63.113	6.134 116.137	1.00 23.44
ATOM	2389		63.108	6.631 117.264	
ATOM	2390		62.530	6.711 115.092	1.00 23.87
ATOM	2391				1.00 22.19
-		_	61.839	7.993 115.216	1.00 26.50
ATOM	2392		61.266	8.416 113.864	1.00 24.16
ATOM	2393		60.715	7.878 116.237	1.00 27.86
ATOM	2394		60.556	8.728 117.117	1.00 22.47
MOTA	-2395		59.940	6.808 116.110	1.00 23.27
ATOM	2396	CA LEU A 302	58.818	6.566 116.996	1.00 26.50
ATOM	2397	CB LEU A 302	58.036	5.354 116.483	1.00 26.02
ATOM	2398	CG LEU A 302	56.866	4.798 117.291	1.00 29.73
ATOM	2399		55.983	3.938 116.394	1.00 31.01
ATOM	2400		57.394	4.001 118.465	
ATOM	2401		59.246	6.373 118.451	1.00 32.99
ATOM	2402	O LEU A 302	58.648	0.3/3 118.451	1.00 27.49
				6.946 119.358	1.00 25.22
ATOM	2403	N ALA A 303	60.289	5.580 118.672	1.00 27.85
MOTA	2404	CA ALA A 303	60.765	5.320 120.024	1.00 27.33
ATOM	2405	CB ALA A 303	61.854	4.269 119.990	1.00 29.17
MOTA	2406	C ALA A 303	61.27 9	6.580 120.714	1.00 26.64
MOTA	2407	O ALA A 303	. 60.944	6.849 121.875	1.00 23.18
ATOM	2408	N ARG A 304	62.092	7.354 120.003	1.00 27.48
MOTA	2409	CA ARG A 304	62.648	8.570 120.581	1.00 25.46
ATOM	2410	CB ARG A 304	63.773	9.136 119.704	1.00 21.31
MOTA	2411	CG ARG A 304	65 .005	8.231 119.562	1.00 25.98
ATOM	2412	CD ARG A 304	66.153	9.042 118.951	1.00 27.87
ATOM	2413	NE ARG A 304	65.647	9.766 117.796	1.00 36.76
ATOM	2414	CZ ARG A 304	66.207	10.838 117.261	1.00 30.79
ATOM	2415	NH1 ARG A 304	67.323	11.345 117.768	
ATOM	2416	NH2 ARG A 304	65.623		1.00 30.11
ATOM	2417	C ARG A 304		11.419 116.225	1.00 36.07
			61.585	9.634 120.803	1.00 25.46
ATOM	2418	O ARG A 304	61.519	10.237 121.876	1.00 24.23
MOTA	2419	N ALA A 305	60.741	9.854 119.802	1.00 22.22
ATOM	2420	CA ALA A 305	59.700	10.868 119.910	1.00 26.70
ATOM	2421	CB ALA A 305	58.914	10.960 118.607	1.00 28.14
MOTA	2422	C ALA A 305	58.749	10.626 121.072	1.00 25.54
ATOM	2423	O ALA A 305	58 513	11.520 121.883	1.00 24.17
MOTA	2424	N TRP A 306	58 189	9.426 121.160	1.00 25.66
MOTA	2425	CA TRP A 306	57.270	9.157 122.253	1.00 28.01
ATOM	2426	CB TRP A 306	56.454	7.873 122.012	1.00 18.66
ATOM	2427	CG TRP A 306	55.382		
ATOM	2428	CD2 TRP A 306	54.709	7.019 120.240	1.00 24.88
ATOM	2429	CE2 TRP A 306	53.725	7.646 119.442	
ATOM	2430	CE3 TRP A 306			1.00 23.98
			54.839	5.623 120.181	1.00 23.26
ATOM	2431	CD1 TRP A 306	54.795	9.228 120.599	1.00 20.24
MOTA	2432	NEI TRP A 306	53.799	8.995 119.681	1.00 24.18
MOTA	2433	CZ2 TRP A 306	52.875	6.926 118.590	1.00 24.30
MOTA	2434	CZ3 TRP A 306	53.993	4.906 119.335	1.00 23.89
MOTA	2435	CH2 TRP A 306	53.024	5.562 118.550	1.00 24.12
MOTA	2436	C TRP A 306	57.969	9.113 123.605	1.00 27.93
MOTA	2437	O TRP A 306	57.330	9.319 124.637	1.00 28.58
ATOM	2438	N THR A 307	59.273	8.851 123.615	1.00 26.76
ATOM	2439	CA THR A 307	60.000	8.850 124.881	1.00 22.81
ATOM	2440	CB THR A 307	61.457	8.319 124.730	1.00 25.54
ATOM	2441	OG1 THR A 307	61.435	6.902 124.504	1.00 23.34
	2442	CG2 THR A 307	62.269	8.599 125.988	
MOTA		COL LINE IS TO.	02.203	0.333 163.300	1.00 24.03

ATOM	2443 C THR A 307		60.027	7 10.28	8 125.396	1 00	26.54
ATOM	2444 O THR A 307		59.925				25.34
ATOM	2445 N LEU A 308		60.152	11.24	7 124.478	1.00	21.65
ATOM	2446 CA LEU A 308		60.172	12.65		1.00	21.41
MOTA	2447 CB LEU A 308		60.442	13.55	B 123.642	1.00	19.20
ATOM	2448 CG LEU A 308		61.797	13.38	5 122.938	1.00	21.31
ATOM	2449 CD1_LEU A 308		61.900		2 121.774	1.00	21.75
MOTA	2450 CD2 LEU A 308		62.937	13.62	2 123.915	1.00	19.26
ATOM	2451 C LEU A 308		58.811			1.00	25.18
MOTA	2452 O LEU A 308		58.731				21.35
ATOM	2453 N ILE A 309	-	57.743				21.74
MOTA	2454 CA ILE A 309 2455 CB ILE A 309		56.394	12.799			19.23
MOTA			55.337				19.63
ATOM	2457 CG1 ILE A 309		53.945			1.00	19.54
ATOM	2458 CD1 ILE A 309		55.403 55.118				20.80
ATOM	2459 C ILE A 309		56.228				20.08
ATOM	2460 O ILE A 309		55.731				23.97
ATOM	2461 N TRP A 310	•	56.652	10.977			21.38 26.45
ATOM	2462 CA TRP A 310		56.525	10.342			28.35
ATOM	2463 CB TRP A 310		56.940	8.872		1 00	23.95
ATOM	2464 CG TRP A 310		56.874	8.203		1 00	29.60
MOTA	2465 CD2 TRP A 310		55.697	7.967		1.00	31.40
ATOM-	2466 CE2 TRP A 310		56.115	7.390		1.00	32.47
ATOM	2467 CE3 TRP A 310		54.329	8.189		1.00	
MOTA	2468 CD1 TRP A 310		57.926	7.770		1.00	33.42
ATOM	2469 NE1 TRP A 310		57.480	7.282		1.00	
ATOM	2470 CZ2 TRP A 310		55.213		132.492	1.00	
ATOM	2471 CZ3 TRP A 310 2472 CH2 TRP A 310		53.432	7.831		1.00	
ATOM ATOM	2472 CH2 IRP A 310 2473 C TRP A 310		53.881	7.259		1.00	
ATOM	2474 O TRP A 310		57.308 56.820		129.293	1.00	
ATOM	2475 N CYS A 311		58.512	11.137 11.535	130.426 128.984	1.00	
ATOM	2476 CA CYS A 311		59.305	12.247		1.00	
ATOM	2477 CB CYS A 311		60.722	12.538		1.00	
ATOM	2478 SG CYS A 311		61.804		129.327	1.00	
ATOM	2479 C CYS A 311		58.612		130.397	1.00	
ATOM	2480 O CYS A 311		58.612		131.570	1.00	28.80
ATOM	2481 N GLU A 312		58.021	14.247		1.00	23.13
MOTA	2482 CA GLU A 312	•	57:308	15.496	129.696	1.00	
ATOM	2483 CB GLU A 312		56.648		128.427	1.00	28.97
ATOM	2484 CG GLU A 312		57.080		127.988	1.00	41.67
ATOM	2485 CD GLU A 312		56.905		129.059	1.00 4	14.21
ATOM	2486 OE1 GLU A 312 2487 OE2 GLU A 312		55.813		129.658	1.00	
ATOM ATOM	2487 OE2 GLU A 312 2488 C GLU A 312		57.860	19.233	129.290	1.00 4	
ATOM	2489 O GLU A 312		56.204 56.120		130.712	1.00 2	
ATOM	2490 N LEU A 313		55.343	15.869	131.751 130.388	1.00 3	30.64
ATOM	2491 CA LEU A 313		54.231		131.266	1.00 3	
ATOM	2492 CB LEU A 313		53.337		130.604	1.00 3	0.21
ATOM	2493 CG LEU A 313		52.493	13.342		1.00 3	14 62
ATOM	2494 CD1 LEU A 313		51.818	12.146		1.00 3	
ATOM	2495 CD2 LEU A 313		51.471	14.357		1.00 2	
ATOM	2496 C LEU A 313		54.685	13.377		1.00 3	
MOTA	2497 O LEU A 313		54.131	13.730		1.00 3	
ATOM	2498 N SER A 314	!	55.688	12.508	132.577	1.00 3	
MOTA	2499 CA SER A 314		56.233	11.880	133.776	1.00 3	
ATOM	2500 CB SER A 314		57.183	10.743	133.388	1.00 3	
MOTA	2501 OG SER A 314		56.517		132.628	1.00 4	
ATOM	2502 C SER A 314		57.002	12.846		1.00 3	
ATOM	2503 O SER A 314		57.339	12.513		1.00 2	10.
MOTA	2504 N GLY A 315		7.312	14.021		1.00 3	
ATOM	2505 CA GLY A 315			14.996		1.00 3	
MOTA	2506 C GLY A 315			14.634		1.00 3	
ATCM	2507 O GLY A 315 2508 N ARG A 316			15.049		1.00 4	
MOTA	TOO N AND M JIO		0.089	13.862	124.181	1.00 3	3.32

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ATOM	2509 CA ARG A 316	61.490	13.501 134.332	2 1 00 30 05
ATOM		61.641	11.982 134.413	
ATOM			11 306 134.413	
ATOM		61.233		1.00 39.11
		61.426		1.00 41.21
ATOM		60.461		1.00 40.97
ATOM		60.524	8.021 134.926	1.00 36.40
MOTA	2515 NH1 ARG A 316	61.511		1.00 38.93
ATOM	2516 NH2 ARG A 316	59.583	7.621 135.768	
ATOM				
ATOM		62.369		
		61.910		1.00 34.27
ATOM		63.633		
ATOM		64.580	14.905 132.619	1.00 44.42
AŤOM	2521 CB GLU A 317	65.901		1.00 46.84
ATOM	2522 CG GLU A 317	65.756	15.996 134.629	
ATOM	2523 CD GLU A 317	65.212	15 112 125 542	
ATOM	2524 OE1 GLU A 317		15.113 135.743	
		65.871	14.101 136.073	
MOTA	2525 OE2 GLU A 317	64.129	15.425 136.287	1.00 67.24
MOTA	2526 C GLU A 317	64.873	13.962 131.462	1.00 38.45
ATOM	2527 O GLU A 317	64.977	12.748 131.636	1.00 38.84
ATOM	2528 N VAL A 318	65.010	14.525 130.275	1.00 37.64
ATOM	2529 CA VAL A 318	65.315	13.720 129.108	
ATOM	2530 CB VAL A 318	64.858	13.720 129.108	
ATOM			14.417 127.810	
		65.192	13.544 126.610	1.00 41.84
ATOM	2532 CG2 VAL A 316	63.364	14.701 127.867	1.00 42.38
MOTA	2533 C VAL A 318	66.822	13.495 129.037	1.00 38.45
MOTA	2534 O VAL A 318	67.598	14.442 128.910	1.00 36.04
ATOM	2535 N PRO A 319	67.261	12.236 129.156	1.00 39.54
ATOM	2536 CD PRO A 319	66.512	10.994 129.397	
ATOM	2537 CA PRO A 319	68.695	10.994 129.397	1.00 40.47
ATOM	2538 CB PRO A 319		11.949 129.088	1.00 43.85
		. 68.745	10.439 129.319	1.00 44.12
ATOM	2539 CG PRO A 319	67.419	9.986 128.745	1.00 46.48
ATOM	2540 C PRO A 319	69.228	12.353 127.718	1.00 43.55
MOTA	2541 O PRO A 319	68.563	12.141 126.708	1.00 43.45
MOTA	2542 N GLU A 320	70.420	12.936 127.689	1.00 42.52
ATOM	2543 CA GLU A 320	71.026	13.380 126.440	
ATOM	2544 CB GLU A 320	72.384		1.00 45.19
ATOM	2545 CG GLU A 320		14.032 126.706	1.00 43.86
		73.121	14.412 125.434	1.00 52.62
ATOM	2546 CD GLU A 320	74.507	14.967 125.697	1.00 52.36
MOTA	2547 OE1 GLU A 320	75.219	15.271 124.720	1.00 56.25
ATOM	. 2548 OE2 GLU A 320	74.883	15.101 126.875	1.00 52.25
ATOM	2549 C GLU A 320	71.223	12.266 125.421	1.00 43.52
ATOM	2550 O GLU A 320	70.876	12.412 124.253	1.00 41.89
ATOM	2551 N LYS A 321	71.781	11.150 125.867	1.00 41.09
ATOM	2552 CA LYS A 321	72.059	10.041 124 060	1.00 43.35
ATOM	2553 CB LYS A 321		10.041 124.969	1.00 43.53
		73.561	9.808 124.879	1.00 42.78
ATOM		74.238	9.340 126.180	1.00 49.38
MOTA	2555 CD LYS A 321	74.272	10.396 127.307	1.00 57.82
ATOM	2556 CE LYS A 321	72.978	10.497 128.129	1.00 53.81
ATOM	2557 NZ LYS A 321	72.660	9.245 128.883	1.00 54.17
ATOM	2558 C LYS A 321	71.407	8.731 125.345	1.00 41.52
ATOM	2559 O LYS A 321	70.954	9 540 125 460	1.00 41.52
ATOM	2560 N LEU A 322		8.540 126.469	1.00 41.98
		71.378	7.820 124.382	1.00 38.64
ATOM	2561 CA LEU A 322	70.815	6.508 124.613	1.00 40.46
MOTA	2562 CB LEU A 322	70.442	5.845 123.289	1.00 42.22
MOTA	2563 CG LEU A 322	69.595	6.632 122.287	1.00 42.92
MOTA	2564 CD1 LEU A 322	69.204	5.737 121.125	1.00 41.13
MOTA	2565 CD2 LEU A 322	68.361	7.148 122.967	7 00 44 44
ATOM	2566 C LEU A 322	71.918		1.00 44.41
	2567 O LEU A 322		5.702 125.268	1.00 41.36
ATOM		73.079	5.825 124.884	1.00 44.16
MOTA	2568 N ASN A 323	71.579	4.894 126.265	1.00 39.89
MOTA	2569 CA ASN A 323	72.594	4.067 126.895	1.00 40.96
MOTA	2570 CB ASN A 323	72.136	3.556 128.259	1.00 43.00
ATOM	2571 CG ASN A 323	70.787	2.886 128.202	1.00 45.59
MOTA	2572 OD1 ASN A 323	70.482	2.151 127.264	1.00 45.71
ATOM	2573 ND2 ASN A 323	69.975	3.114 129.224	
			J.114 147.444	1.00 48.08
ATOM	2574 C ASN A 323	72.828	2.894 125.954	1.00 44.88

•	
ATOM 2575 O ASN A	322 70 404
1.2	4.737 124.333 1 (1) 46 A
- 1017	74.809 7.062.126.260 1.00 1.00
ATOM 2577 CA ASN A	344 74 172 0 030 135 404
ATOM 2578 CB ASN A	324 25 25 25 25 25 25 25 25 25 25 25 25 25
ATOM 2579 CG ASN A	
	75.960 -0.711 124.888 1.00 60 0
ATOM 2580 OD1 ASN A	324 75 344 -1 702 104
ATOM 2581 ND2 ASN A	324 1.00 66.9
ATOM 2582 C ASN A	
TION A	72.979 -0.070 125 267 1 00 47 1
ATOM 2583 O ASN A	1324 72.784 -0 644 124 107 7 7
ATOM 2584 N LYS A	725 72 73 73 74 75 75 75 75 75 75 75 75 75 75 75 75 75
ATOM 2585 CA LYS A	
	325 71.106 -1 221 126 220 1 22
ATOM 2586 CB LYS A	345 70.428 -1 329 127 605
ATOM 2587 CG LYS A	375 7.020 127.093 1.00 47.65
1.	71.292 -1.83/ 128.858 1 00 54 21
	325 72.160 -0.750 129 526 1 00 FC 07
ATOM 2589 CE LYS A	775
ATOM 2590 NZ LYS A	7.25 120.071 1.00 57.45
	74.091 0.816 129.307 1.00 58 33
	345 70.062 -0.791 125 206 1 20
ATOM 2592 O LYS A	775 (0 4m.
ATOM 2593 N ALA A	336
	09.832 0.514 125.213 1.00 41 43
	346 68 861 1 054 124 226 1 22
ATOM 2595 CB ALA A	226
ATOM 2596 C ALA A	
	09.365 0.940 122.838 1 00 43 90
	346 68.595 0.625 121 026 1 02
ATOM 2598 N LYS A	
ATOM 2599 CA LYS A	227 1.00 43.46
DID A	71.233 1.120 121.296 1.00 43 26
11 ,	72.723 1.484 121 311 1 00 44 77
ATOM 2601 CG LYS A	177 77 77 77
ATOM 2602 CD LYS A	222 121.800 1.00 50.87
ATOM 2603 CE LYS A	3.200 221,747 1.111 57 77
OID A	12' 14'310 4'508 122'322 1 00 E1 30
ATOM 2604 NZ LYS A 3	4/ 74.256 5.670 121 715 1 46
ATOM 2605 C LYS A 3	27 2.00 32.72
ATOM 2606 O LYS A 3	27 1.00 41.29
ATOM 2607 N GLU A 3	2' '0.625 -0.437 119.592 1.00 38 83
	28 71.403 -1.278 121 526 1 00 20 00
ATOM 2608 CA GLU A 3	
ATOM 2609 CB GLU A 3	2.000 121.090 1.00 42.40
- 020 A J	40 /3.164 m3 /32 122 20E 1 00 1 m
ATOM 2611 CD GLU A 3	28 74 096 -3 520 120 002 1 22
ATOM 2612 OE1 GLU A 3	2.025 120.965 1.00 52.11
ATOM 2613 OE2 GLU A 3	100 55.44
ATOM 2614 C GLU A 3	74.785 -2.558 120.584 1.00 51.64
	48 69.825 -3 030 120 010 1 00 -2
ATOM 2615 O GLU A 3:	08 60 536 3 545
ATOM 2616 N LEU A 32	3.042 113.939 1.00 37.35
	_ ~~~~ ~~~~ ~~~~~~~~~~~~~~~~~~~~~~~~~~~
	9^{-9} 9^{-1} , 499 -2.717 121.380 1 00 36 03
ATOM 2618 CB LEU A 32	9 66.646 -1 959 122 400 1 00 74
ATOM 2619 CG LEU A 32	1.00 34.66
ATOM 2620 CD1 LEU A 32	2.22.213 1.00 33.88
	3 . 04./55 -3.572 122.351 1.00 36 21
1. 32	9 64 391 -1 269 122 240 2 00 2
ATOM 2622 C LEU A 32	9 67.120 -2 269 119 071 1 00 7
ATOM 2623 O LEU A 32	1.00 33.64
ATOM 2624 N LEU A 33	
Anon 2024 N LEU A 33	0/.333 -0.990 119 681 1 nn 22 20
ATOM 2625 CA LEU A 33	0 67.004 -0 461 118 366 1 00 36 75
ATOM 2626 CB LEU A 33	2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2
ATOM 2627 CG LEU A 330	
	, 60.514 1.958 119.505 1 00 31 51
ATOM 2628 CD1 LEU A 330	56.857 3 404 119 904 1 99 99
ATOM 2629 CD2 LEU A 330	5. 100 22.11
ATOM 2630 C LEU A 330	57 75 1.00 29.69
1, 550	07.729 -1.201 117.246 1.00 36.49
550	67.142 -1 493 116 210 1 00 25 61
ATOM 2632 N LYS A 331	CO 00-
ATCM 2633 CA LYS A 331	60 305 -1.303 117.435 1.00 37.63
200 11 001	09.786 -2.205 116.446 1.00 41 32
	1.456 -2.272 116.874 7 00 44 74
ATOM 2635 CG LYS A 331	71 00
ATOM 2636 CD LYS A 331	7,7,7,100
	74 7
	$\frac{4.315}{100}$ -1.889 116.765 1 no.52 71
ATCM 2638 NZ LYS A 331	77 000
ATOM 2639 C LYS A 331	60 250
ATOM 2640 0 LYS A 331	50 7.0 42.33
Prov. 2040 O PAR W 331	69.310 -4.086 115.042 1.00 42.68
	1.00 42.00

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MOTA	2643	N SER A 33	68.734	-4.270 117.200	1.00 41.56
ATOM	2642		68.226	-5.629 117.039	1.00 46.88
MOTA	2643			-6.298 118.400	1.00 42.19
ATOM	2644		-	-5.714 119.096	1.00 39.55
MOTA	2645			-5.687 116.297	1.00 48.58
MOTA	2646			-6.774 116.017	1.00 45.78
MOTA	2647			-4.531 115.979	1.00 48.27
ATOM	2648			-4.503 115.292	1.00 51.82
MOTA	2649 2650			-3.119 115.402	1.00 52.16
MOTA MOTA	2651			-3.122 114.683	1.00 52.64
ATOM	2652			-2.765 116.871 -1.402 117.077	1.00 52.70
ATOM	2653			-4.887 113.820	1.00 56.70 1.00 53.43
ATOM	2654			~4.675 113.145	1.00 56.45
ATOM	2655			-5.461 113.344	1.00 55.53
ATOM	2656			-5.892 111.962	1.00 59.58
ATOM	2657	CB ASP A 334	4 62.845	-7.040 111.918	1.00 62.69
ATOM	2658	CG ASP A 334	4 61.546	-6.712 112.664	1.00 66.23
ATOM	2659	OD1 ASP A 334		-5.814 112.227	1.00 63.25
ATOM	2660	OD2 ASP A 334		-7.354 113.704	1.00 63.45
ATOM	2661	C ASP A 334		-4.705 111.125	1.00 60.81
ATOM	2662	O ASP A 334		-4.673 110.681	1.00 59.47
ATOM	2663	N PHE A 335 CA PHE A 335		-3.736 110.889	1.00 60.00
ATOM	2664 2665	CA PHE A 335 CB PHE A 335		-2.545 110.147	1.00 59.37
MOTA MOTA	2666	CG PHE A 335		-1.298 110.952 -0.013 110.275	1.00 53.38
ATOM	2667	CD1 PHE A 335		0.189 109.770	1.00 44.32
ATOM	2668	CD2 PHE A 335		0.993 110.127	1.00 49.91
MOTA	2669	CE1 PHE A 335		1.373 109.122	1.00 42.74
ATOM	2670	CE2 PHE A 335	64.526	2.180 109.483	1.00 46.40
ATOM	2671	CZ PHE A 335		2.370 108.978	1.00 42.77
ATOM	2672	C PHE A 335		-2.399 108.696	1.00 60.85
MOTA	2673	O PHE A 335		-2.914 107.785	1.00 66.28
ATOM	2674	N GLU A 336		-1:671 108.493	1.00 57.40
MOTA	2675 2676	CA GLU A 336 CB GLU A 336		-1.411 107.174 -2.579 106.211	1.00 58.96
MOTA MOTA	2677	CG GLU A 336		-2.579 106.211 -2.377 104.846	1.00 62.66 1.00 68.51
ATOM	2678	CD GLU A 336		-3.590 103.943	1.00 73.21
ATOM	2679	OE1 GLU A 336		-4.678 104.333	1.00 73.30
ATOM	2680	OE2 GLU A 336		-3.457 102.843	1.00 75.74
MOTA	2681	C GLU A 336	65.460	-0.124 106.576	1.00 55.70
ATOM	2682	0 GLU A 336	64.281	-0.023 106.253	1.00 55.28
MOTA	2683	N GLU A 337	66.338	0.857 106.432	1.00 54.75
ATOM	2684	CA GLU A 337	65.986	2.167 105.905	1.00 55.99
ATOM	2685	CB GLU A 337	67.221	3.065 105.983	1.00 51.75
ATOM	2686 2687	CG GLU A 337 CD GLU A 337	66.926 66.184	4.536 106.092 4.891 107.366	3 00 52.28.
MOTA MOTA	2688	OE1 GLU A 337	66.705	4.640 108.474	:.00 43.72 : 00 42.21
ATOM	2689	OE2 GLU A 337	65.072	5.425 107.256	1.00 47.31
ATOM	2690	C GLU A 337	65.485	2.064 104.460	1.00 57.56
ATOM	2691	O GLU A 337	66.087	1.377 103.639	1.00 58.29
ATOM	2692	N PHE A 338	64.385	2.745 104.151	1.00 60.26
ATOM	2693	CA PHE A 338	63.814	2.710 102.805	1.00 61.69
ATOM	2694	CB PHE A 338	62.561	3.582 102.723	1.00 60.86
MOTA	2695	CG PHE A 338	61.845	3.493 101.401	1.00 61.92
ATOM	2696	CD1 PHE A 338	61.054	2.391 101.094	1.00 62.99
ATOM	2697	CD2 PHE A 338	61.970	4.508 100.458	1.00 62.74
MOTA	2698	CE1 PHE A 338 CE2 PHE A 338	60.392 61.315	2.302 99.868	1.00 65.11
ATOM	2699	CE2 PHE A 338 CZ PHE A 338	60.523	4.428 99.228 3.322 98.934	1.00 64.62
ATOM	2700 2701	C PHE A 338	64.818	3.322 98.934 3.208 101.773	1.00 63.87 1.00 64.33
ATOM ATOM	2702	O PHE A 338	64.803	2.781 100.616	1.00 62.45
ATOM	2702	N ASP A 339	65.677	4.130 102.194	1.00 64.11
ATOM	2704	CA ASP A 339	66.689	4.684 101.310	1.00 67.42
ATCM	2705	CB ASP A 339	66.565	6.206 101.248	1.00 67.20
ATOM	2706	CG ASP A 339	67.647	6.838 100.402	1.00 68.03

ATOM	2707 OD1 ASP A 339	67.796	6.432 99.22	1 00 72 20
ATOM		68.346		
ATOM		68.088		
ATOM				
		68.511		
ATOM		68.796		
ATOM	<u> </u>	70.151	3.111 101.149	1.00 73.59
ATOM	2713 CB ASP A 340	70,778		3 1.00 75.47
MOTA	2714 CG ASP A 340	69.953	1.511 99.195	
ATOM	2715 OD1 ASP A 340	69.761	0.449 99.823	1.00 76.03
ATOM	2716 OD2 ASP A 340	69.492		
MOTA	2717 C ASP A 340	09.492	1.718 98.054	
		- 71.069	4.155 101.766	
ATOM	2718 O ASP A 340	71.618	3.946 102.845	1.00 73.08
MOTA	2719 N GLU A 341	71.242	5.275 101.074	1.00 73.15
ATOM	2720 CA GLU A 341	72.112	6.341 101.557	1.00 74.56
AŢOM	2721 CB GLU A 341	72.917	6.924 100.390	1.00 77.06
ATOM	2722 CG GLU A 341	73.878	8.034 100.792	
ATOM	2723 CD GLU A 341	74.924	7 573 101 702	
ATOM	2724 071 071 0 244		7.571 101.794	
	2725 OE2 GLU A 341		6.669 101.450	
MOTA		74.951	8.106 102.924	
ATOM	2726 C GLU A 341	71.327	7.453 102.245	1.00 72.47
MOTA	. 2727 O GLU A 341	70.822	8.364 101.589	1.00 76.75
MOTA	2728 N VAL A 342	71.228	7.381 103.566	1.00 67.86
MOTA	2729 CA VAL A 342	70.503	8.393 104.323	1.00 64.84
ATOM	2730 CB VAL A 342	69.160		
ATOM	2731 CG1 VAL A 342	68.256		
MOTA	2732 CG2 VAL A 342		7.494 103.701	1.00 67.95
ATOM	2733 C VAL A 342	69.400	6.637 105.722	1.00 65.37
		71.305	8.871 105.520	1.00 61.75
MOTA		71.375	10.066 105.795	1.00 64.14
ATOM	2735 N ASP A 343	71.912	7.925 106.225	1.00 56.79
ATOM-	2736 CA ASP A 343	72.692	8.229 107.417	1.00 54.53
ATOM	2737 CB ASP A 343	73.707	9.340 107.158	1.00 56.31
ATOM	2738 CG ASP A 343	74.531	9.660 108.388	1.00 58.81
MOTA	2739 OD1 ASP A 343	75.298	10.644 108.357	1.00 50.01
ATOM	2740 OD2 ASP A 343	74.420		1.00 65.36
ATOM	2741 C ASP A 343		8.918 109.387	1.00 54.29
ATOM	2742 O ASP A 343	71.765	8.675 108.534	1.00 50.70
		71.442	9.859 108.651	1.00 46.00
ATOM		71.328	7.717 109.341	1.00 46.20
ATOM	2744 CA ARG A 344	70.452	8.004 110.463	1.00 41.18
MOTA	2745 CB ARG A 344	69.121	7.268 110.299	1.00 39.81
ATOM	2746 CG ARG A 344	68.289	7.711 109.098	1.00 35.08
ATOM	2747 CD ARG A 344	68.036	9.211 109.121	1.00 28.37
ATOM	2748 NE ARG A 344	67.157	9.645 108.036	1.00 30.90
ATOM	2749 CZ ARG A 344	67.013	10.909 107.649	1.00 30.90
ATOM	2750 NH1 ARG A 344	67.693	11.874 108.258	1.00 31.05
ATOM	2751 NH2 ARG A 344	66.201		1.00 30.49
ATOM	2752 C ARG A 344		11.212 106.646	1.00 31.76
		71.147	7.561 111,742	1.00 38.46
ATOM		70.516	7.370 112.773	1.00 34.99
ATOM	2754 N SER A 345	72.464	7.418 111.662	1.00 33.97
MOTA	2755 CA SER A 345	73.261	6.981 112.795	1.00 33.68
MOTA	2756 CB SER A 345	74.742	6.972 112.404	1.00 39.11
MOTA	2757 OG SER A 345	75.163	8.260 111.990	1.00 42.80
ATOM	2758 C SER A 345	73.054	7.826 114.053	
ATOM	2759 O SER A 345	73.100	7.314 115.167	1.00 31.83
ATOM	2760 N TYR A 346	72.819	0 110 111 077	1.00 24.35
			9.119 113.877	1.00 33.10
ATOM			10.003 115.015	1.00 34.50
ATOM	2762 CB TYR A 346		11.437 114.522	1.00 35.16
MOTA	2763 CG TYR A 346		11.615 113.659	1.00 39.69
ATOM	2764 CD1 TYR A 346	69.909	11.814 114.227	1.00 36.57
MOTA	2765 CE1 TYR A 346		11.940 113.424	1.00 40.23
ATOM	2766 CD2 TYR A 346		11.544 112.270	1.00 39.04
MOTA	2767 CE2 TYR A 346	70.131	11.667 111.463	1 00 30 66
MOTA	2768 CZ TYR A 346	68.890		1.00 38.65
	2769 OH TYR A 346		11.864 112.041	1.00 37.64
MOTA			11.982 111.234	1.00 32.48
MOTA	2770 C TYR A 346	71.432	9.560 115.874	1.00 37.72
ATCM	2771 O TYR A 346.	71.396	9.829 117.074	1.00 35.48
ATCM	2772 N MET A 347	70.472	8.869 115.265	1.00 35.36
				· · ·

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ATOM	2773	3 CA MET A 347	69.295	8.418 116.004	1.00 36.97
MOTA	2774		68.226		1.00 33.45
MOTA	2775	G MET A 347	67.853		1.00 28.09
ATOM	2776	SD MET A 347	66.471		
MOTA	2777		67.058		
ATOM-	2778	3 C MET A 347	69.632		1.00 34.67
MOTA	2779		68.890		1.00 33.46
MOTA	2780		70.74 7	6.663 116.880	1.00 36.42
MOTA	2781		71.137	5.641 117.846	1.00 34.57
MOTA	2782		71.841	4.476 117.152	1.00 34.16
MOTA	2783		71.066		1.00 35.61
ATOM	2784		72.010		1.00 36.58
MOTA	2785		69.906		
MOTA	2786		72.075	6.232 118.885	1.00 37.10
MOTA	2787		72.583	5.513 119.745	1.00 37.75
MOTA	2788 2789		72.295	7.541 118.801	1.00 38.31
MOTA	2790		. 73.192	8.231 119.726	1.00 42.86
ATOM ATOM	2791	CB GLU A 349 . CG GLU A 349	74.150 74.942	9.136 118.948	1.00 47.04
ATOM	2792	CD GLU A 349	75.828	8.426 117.867	1.00 53.27
ATOM	2793	OE1 GLU A 349	76.681	7.325 118.410 7.619 119.279	1.00 58.36
ATOM	2794	OE2 GLU A 349	75.669	6.166 117.963	1.00 59.59
ATOM	2795	C GLU A 349	72.458	9.080 120.752	1.00 59.17
ATOM	2796	O GLU A 349	72.564		1.00 41.72 1.00 42.27
ATOM	2797	N THR A 350	71.716	10.067 120.261	1.00 42.27
ATOM	2798	CA THR A 350	70.992	10.976 121.135	1.00 40.78
ATOM	2799	CB THR A 350	71.468	12.418 120.900	1.00 41.69
MOTA	2800	OG1 THR A 350	71.359	12.733 119.508	1.00 43.25
MOTA	2801	CG2 THR A 350	72.918	12.575 121.325	1.00 44.11
ATOM	2802	C THR A 350	. 69.474	10.911 120.968	1.00 41.47
MOTA	2803	O THR A 350	68.968	10.608 119.884	1.00 39.72
ATOM	2804	N LEU A 351	68.760	11.199 122.054	1.00 38.68
ATOM	2805	CA LEU A 351	67.299	11.185 122.056	1.00 35.91
ATOM	2806	CB LEU A 351	66.763	11.210 123.487	1.00 35.94
ATOM	2807 2808	CG LEU A 351	66.752	9.890 124.251	1.00 37.67
ATOM ATOM	2809	CD1 LEU A 351 CD2 LEU A 351	66.290	10.118 125.677	1.00 40.11
ATOM	2810	C LEU A 351	65.813 66.679	8.920 123.544	1.00 39.29
ATOM	2811	O LEU A 351	65.747	12.342 121.294 12.149 120.512	1.00 37.76
ATOM	2812	N LYS A 352	67.192	13.544 121.525	1.00 34.86 1.00 34.79
ATOM	2813	CA LYS A 352	66.651	14.724 120.870	1.00 34.79
ATOM	2814	CB LYS A 352	66.676	15.911 121.835	1.00 36.48
MOTA	2815	CG LYS A 352	66.062	15.580 123.179	1.00 42.08
MOTA	2816	CD LYS A 352	66.202	16.701 124.196	1.00 43.22
ATOM	2817	CE LYS A 352	65.349	17.901 123.845	1.00 49.81
ATOM	2818	NZ LYS A 352	65.342	18.880 124.972	1.00 52.70
ATOM	2819	C LYS A 352	67.425	15.063 119.610	1.00 38.77
MOTA	2820	O LYS A 352	68.654	15.098 119.601	1.00 36.21
ATOM	2821	N ASP A 353	66.697	15.293 118.530	1.00 37.69
MOTA	2822	CA ASP A 353	67.337	15.647 117.286	1.00 39.89
ATOM	2823	CB ASP A 353	66.532	15.075 116.110	
MOTA	2824	CG ASP A 353	65.058	15.368 116.211	1.00 47.99
MOTA	2825 2826	OD1 ASP A 353 OD2 ASP A 353	64.253	14.623 115.593	1.00 38.06
MOTA	2827	C ASP A 353	64.706	16.352 116.898	1.00 52.94
ATOM	2828	O ASP A 353	67.457	17.165 117.247	1.00 39.20
MOTA MOTA	2829	N PRO A 354	66.890 68.244	17.861 118.092 17.696 116.302	1.00 35.66
ATOM	2830	CD PRO A 354	69.047	17.696 116.302 17.005 115.279	1.00 40.75 1.00 40.05
ATOM	2831	CA PRO A 354	68.426	19.145 116.179	1.00 40.05
ATOM	2832	CB PRO A 354	69.534	19.250 115.140	1.00 36.24
ATOM	2833	CG PRO A 354	69.190	18.095 114.225	1.00 39.41
ATOM	2834	C PRO A 354	67.144	19.780 115.689	1.00 37.28
MOTA	2835	O PRO A 354	66.299	19.106 115.094	1.00 31.87
ATOM	2836	N TRP A 355	66.993	21.074 115.934	1.00 37.87
MOTA	2837	CA TRP A 355	65.804	21.757 115.472	1.00 40.04
MOTA	2838	CB TRP A 355	65.714	23.157 116.080	1.00 42.85

ATOM	2839	CG TRP A 355	64.333	3 23.471 116.569	1.00 53.01
ATOM	2840				
			63.439		
MOTA	2841		62.234		1.00.52.59
MOTA	2842	CE3 TRP A 355	63.538	25.422 115.028	1.00.52.82
MOTA	2843	CD1 TRP A 355	63.655		
	2844				
MOTA			62.393		
ATOM	2845	CZ2 TRP A 355	61.134	25.197 116.532	1.00 51.28
MOTA	2846	CZ3 TRP A 355	62.444	26.245 114.779	1.00 53.22
ATOM	2847		61.257	-	
					1.00 50.50
MOTA	2848		65.935	21.836 113.954	1.00 37.34
ATOM	2849	O TRP A 355	67.041	21.929 113.422	1.00 39.01
ATOM	2850	N ARG A 356	64.809		1.00 36.31
	2851	CA ARG A 356			
ATOM			64.797		1.00 35.22
MOTA	2852	CB ARG A 356	64.317	20.469 111.252	1.00 33.35
MOTA	2853	CG ARG A 356	65.310	19.340 111.564	1.00 34.50
ATOM	2854	CD ARG A 356	64.729		
		_			1.00 28.06
MOTA	2855	NE ARG A 356	. 65.745	16.956 111.870	1.00 24.79
MOTA	2856	CZ ARG A 356	65.499	15.703 112.236	1.00 27.56
MOTA	2857	NH1 ARG A 356	64.253	15.237 112.259	1.00 19.09
	2858	NH2 ARG A 356	66.502	· · · · · · · · · · · · · · · · · · ·	
ATOM				14.918 112.604	1.00 21.34
ATOM	285,9	C ARG A 356	63.874	22.955 111.390	1.00 33.74
ATOM	2860	O ARG A 356	62.746	22.732 110.950	1.00 34.68
MOTA	2861	N GLY A 357	64.361		1.00 35.60
ATOM	2862	CA GLY A 357			
		_	63.556	25.345 111.220	1.00 35.43
ATOM	2863	C GLY A 357	63.719	25.932 109.830	1.00 38.08
ATOM	2864	O GLY A 357	64.112	25.250 108.885	1.00 37.29
ATOM	2865	N GLY A 358	63.406	27.218 109.721	1.00 39.67
	2866	CA GLY A 358	63.493		
ATOM				27.925 108.457	1.00 36.36
MOTA	2867	C GLY A 358	62.398	28.966 108.499	1.00 39.45
ATOM	2868	O GLY A 358	61.763	29.131 109.539	1.00 37.58
ATOM	2869	N GLU A 359	62.163	29.662 107.391	1.00 40.89
	2870	-	61.121		
MOTA				30.682 107.358	1.00 41.37
MOTA	2871	CB GLU A 359	61.310	31.627 106.172	1.00 44.64
ATOM	2872	CG GLU A 359	60.956	30.977 104.848	1.00 52.13
ATOM	. 2873	CD GLU A 359	60.833	31.973 103.708	1.00 59.14
	2874	OE1 GLU A 359		_	
MOTA			60.448	31.551 102.593	1.00 60.47
ATOM	2875	OE2 GLU A 359	61.119	33.173 103.923	1.00 57.77
MOTA	2876	C GLU A 359	59.770	30.006 107.200	1.00 38.02
ATOM	2877	O GLU A 359	59.689	28.828 106.850	1.00 35.29
MOTA	. 2878		58.708	30.762 107.441	1.00 36.81
MOTA	2879	CA VAL A 360	57.363	30.237 107.291	1.00 35.97
ATOM	2880	CB VAL A 360	56.401	30.789 108.368	1.00 34.90
ATOM	2881	CG1 VAL A 360	54.999	30.251 108.133	1.00 36.53
	2882	CG2 VAL A 360			
ATOM			56.888	30.393 109.755	1.00 37.06
ATOM	2883	C VAL A 360	56.886	30.690 105.928	1.00 36.74
ATOM	2884	O . VAT V 300	56.661	31.881 105.712	1.00 34.90
ATOM	2885	N ARG A 361	56.753	29.741 105.004	1.00 35.48
ATOM	2886	CA ARG A 361	56.301	30.049 103.652	1.00 38.21
ATOM	2887	CB ARG A 361	56.152	28.776 102.815	1.00 39.76
ATCM	2888	CG ARG A 361	57.416	28.342 102.098	1.00 39.93
	2889	CD ARG A 361	57.225	26.963 101.486	
ATOM					1.00 38.68
ATCM	2890	NE ARG A 361	57.112	25.940 102.525	1.00 39.72
ATOM	2891	CZ ARG A 361	56.952	24.643 102.286	1.00 38.79
MOTA	2892	NH1 ARG A 361	56.881	24.200 101.036	1.00 32.40
ATOM	2893	NH2 ARG A 361	56.899	23.785 103.297	1.00 36.58
ATOM	2894	C ARG A 361	54.996	30.807 103.603	1.00~38.98
ATOM	2895	O _ ARG A 361	54.120	30.636 104.452	1.00 39.07
	3896	N LYS A 362	54.880	31.634 102.573	1.00 39.95
ATCM					
MOTA	2897	CA LYS A 362	53.709	32.459 102.339	1.00 42.73
ATCM	2898	CB LYS A 362	53.931	33.301 101.078	1.00 44.92
ATCM	2899	CG LYS A 362	54.995	34.390 101.219	1.00 55.45
		CD LYS A 362	56.351	33.842 101.671	1.00 58.28
ATOM					
ATCM		CE LYS A 362	56.907	32.809 100.697	1.00 57.04
ATOM	2902	NZ LYS A 362	58.224	32.283 101.151	1.00 58.75
ATOM		C LYS A 362	52.434	31.634 102.200	1.00 40.51
			51.391	31.996 102.748	1.00 36.10
ATCM	2904	O LYS A 362	22.221	34.330 402.748	1.00 30.10

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Figure 18-45

ATO	M 2905 N GLU A 363	52.506 30.527 101.469 1.00 37.79
OTA	M 2906 CA GLU A 363	
ATO	M 2907 CB GLU A 363	
ATO	M 2908 CG GLU A 363	52 500
ATO	M 2909 CD GLU A 363	
ATO]		
ATO	M 2911 OE2 GLU A 363	54 446 43.03
ATO	M 2912 C GLU A 363	2.00 40.44
ATO	M 2913 O GLU A.363	
ATO		2.00 34.19
ATON		
ATON	11.04	
ATON	11 503	
ATON	2918 CG2 VAL A 364	52.018 27.676 107.160 1.00 36.90
ATOM	2919 C VAL A 364	53.198 26.867 105.109 1.00 29.56 50.506 29.574 105.589 1.00 34 33
ATOM	1 2920 O VAL A 364	
ATOM		
ATOM	505	51.027 30.797 105.499 1.00 38.12
ATOM	2923 CB LYS A 365	50.381 31.952 106.119 1.00 36.77
ATOM		51.255 33.204 105.969 1.00 37.98
ATOM		52.629 33.084 106.610 1.00 37.99
ATOM	505	53.449 34.357 106.429 1.00 35.50
ATOM	2927 NZ LYS A 365	54.837 34.190 107.032 1.00 40.35
ATOM	2928 C LYS A 365	55.674 35.407 106.877 1.00 43.74
ATOM	11 303	49.025 32.191 105.468 1.00 36.62
ATOM		48.038 32.469 106.148 1.00 33.53
ATOM	2931 CA ASP A 366	48.968 32.073 104.147 1.00 37.05
ATOM	2932 CB ASP A 366	47.708 32.278 103.449 1.00 37.72
ATOM	2933 CG ASP A 366	47.906 32.237 101.929 1.00 40.57
ATOM	2934 OD1 ASP A 366	48.833 33.334 101.427 1.00 43.98
ATOM	2935 OD2 ASP A 366	49.078 34.304 102.176 1.00 38.51
ATOM	2936 C ASP A 366	49.297 33.235 100.269 1.00 41.96
ATOM	2937 O ASP A 366	46.670 31.238 103.862 1.00 39.24
ATOM	2938 N THR A 367	45.497 31.562 104.029 1.00 39.04
ATOM	2939 CA THR A 367	47.096 29.990 104.031 1.00 38.99
ATOM	2940 CB THR A 367	46.167 28.935 104.432 1.00 36.80
ATOM	2941 OG1 THR A 367	46.868 27.560 104.527 1.00 33.84
ATOM	2942 CG2 THR A 367	47.332 27.167 103.232 1.00 34.92
ATOM	2943 C THR A 367	45.904 26.509 105.046 1.00 35.11 45.532 29.257 105.786 1.00 36.58
ATOM	2944 O THR A 367	2.00 30,38
ATOM	2945 N LEU A 368	
ATOM	2946 CA LEU A 368	
ATOM	2947 CB LEU A 368	2,00 33,40
ATOM	2948 CG LEU A 368	2.00 54.05
ATOM	2949 CD1 LEU A 368	
MOTA	2950 CD2 LEU A 368	10 37.33
ATOM	2951 C LEU A 368	
ATOM	2952 O LEU A 368	1.00 30.12
ATOM	2953 N GLU A 369	
ATOM	2954 CA GLU A 369	
ATOM	2955 CB GLU A 369	45 100
ATOM	2956 CG GLU A 369	
ATOM	2957 CD GLU A 369	
ATOM	2958 OE1 GLU A 369	15 010 33.31
ATOM	2959 OE2 GLU A 369	2.00 00.00
ATOM	2960 C GLU A 369	2.00 05.05
ATOM	2961 O GLU A 369	2.00 45.25
ATOM	2962 N LYS A 370	42,130 33.283 107.095 1.00 44.62
ATOM	2963 CA LYS A 370	42.983 32.057 105.417 1.00 40.34
ATOM	2964 CB LYS A 370	41.666 31.631 104.977 1.00 43.36
ATOM	2965 CG LYS A 370	41.738 30.773 103.704 1.00 44.79
ATOM	2966 CD LYS A 370	42.032 31.546 102.419 1.00 48.93
ATOM	2967 CE LYS A 370	43.503 31.514 102.019 1.00 51.41
ATOM	2968 NZ LYS A 370	43.921 30.116 101.561 1.00 51.14
ATOM	2969 C LYS A 370	45.339 30.062 101.091 1.00 50.34 40.959 30.848 106.069 1.00 43.09
ATOM	2970 O LYS A 370	20 745 106
ALVM		39.745 30.977 106.248 1.00 41.34

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ATOM	2971 N ALA	. 271	43 315	20.00-		
ATOM		371	41.715		106.802	
		371	41.120		107.86	
ATOM		A 371	42.193		108.594	1.00 39.11
ATOM		371	40.365		108.837	1.00 46.10
ATOM			39.230	29.829	109.210	1.00 46.07
MOTA	2976 N LYS 3		40.981	31.239		1.00 46.62
MOTA	2977 CA LYS 2	372	40.391	32.178		
MOTA	2978 C LYS 2	372	39.052	32.698		1.00 51.67
ATOM	2979 O LYS A		38.294	33.318		1:00 51.6/
ATOM	2980 CB LYS A		41.334			
MOTA	2981 CG LYS A		42.804	33.364		
MOTA	2982 CD LYS A				110.510	
MOTA			43.746	34.131		
			45.216		110.849	
ATOM	2984 NZ LYS A		46.121	34.850	111.081	1.00, 20.00
ATOM	2985 N ALA A		38.751	32.476	108.397	
ATOM	2986 CA ALA A		37.492	32.933	107.806	
MOTA	2987 CB ALA A	373	37.758	33.632	106.480	
MOTA	2988 C ALA A	373	36.524	31.773	107.594	1.00 59.58
ATOM	2989 O ALA A	373	35.432		108.205	1.00 60.89
ATOM	2990. OXT ALA A		36.870		106.822	1.00 60.10
ATOM	3014 CB ALA B	2	54.881	-4.431	56.836	
ATOM	3015 C ALA B	2	53.960	-2.137		1.00 55.77
ATOM	3016 O ALA B	2			56.480	1.00 57.58
ATOM	3017 N ALA B	2	54.920	-1.720	57.131	1.00 56.75
			54.263	-3.672	54.557	1.00 58.22
MOTA		2	53.914	-3.584	56.008	1.00 58.47
MOTA	3019 N LYS B	3	52.919	-1.376	56.151	1.00 52.79
MOTA	3020 CA LYS B	3	52.855	0.022	56.543	1.00 49.68
MOTA	3021 CB LYS B	3	51.643	0.700	55.896	1.00 53.14
ATOM	3022 CG LYS B	3	51.751	0.785	54.377	1.00 53.37
ATOM	3023 CD LYS B	3	50.685	1.681	53.786	1.00 55.40
MOTA	3024 CE LYS B	3	50.808	1.783	52.277	1.00 59.51
MOTA	3025 NZ LYS B	3	52.140	2.323	51.884	1.00 56.88
MOTA	3026 C LYS B	3	52.849	0.238	58:059	1.00 46.83
ATOM	3027 O LYS B	3.	52.389	-0.607	58.830	
ATOM	3028 N VAL B	. 4	53.376	1.385	58.467	1.00 41.63
ATOM	3029 CA VAL B	4	53.483	1.751		1.00 41.46
ATOM	3030 CE VAL B	4	54.893		59.871	1.00 40.85
ATOM	3031 CG1 VAL B	4	55.070	2.288	60.163	1.00 39.55
ATOM	3032 CG2 VAL B	4		2.541	61.648	1.00 41.23
ATOM	3033 C VAL B		55.916	1.306	59.652	1.00 38.96
ATOM	3034 O VAL B	4	52.451	2.813	60.230	1.00 38.92
		4	52.472	3.916	59.691	1.00 42.80
ATOM		5	51.559	2.479	61.157	1.00 34.90
ATOM	3036 CA LYS B	5	50.501	3.396	61.558	1.00 31.22
MOTA	3037 CE LYS B	5	49.133	2.796	61.215	1.00 33.76
ATOM	3038 CG LYS B	5	48.841	2.623	59.726	1.00 36.60
ATOM	3039 CD LYS B	5	48.667	3.964	59.032	1.00 41.48
MOTA	3040 CE LYS B	5	48.234	3.803	57.577	1.00.43.62
MOTA	3041 NZ LYS B	5	49.215	3.025	56.781	1:00 42.53
ATOM	3042 C LYS B	5.	50.512	3.749	63.038	1.00 32.67
ATOM	3043 O LYS B	5	51.012	2.995	63.878	1.00 25.78
ATOM	3044 N LEU B	6	49.937	4.906		
ATOM	3045 CA LEU B	6	49.821		63.343	1.00 27.07 1.00 31.09
ATOM	3046 CB LEU B	6		5.379	64.712	1.00 31.09
			50.596	6.696	64.896	1.00 30.13
ATOM		6	50.691	7.340	66.285	1.00 28.09
ATOM	3048 CD1 LEU B	6	49.333	7.827	66.728	1.00 38.87
ATOM	3049 CD2 LEU B	6	51.248	6.338	67.282	1.00 24.87
MOTA	3050 C LEU B	6	48.324	5.594	64.924	1.00-29.52
MOTA	3051 0 LEU 9	6	47.669		64.149	1.00 33.36
MOTA	3052 N ILE B	7	47.777		65.960	1.00 28.02
MOTA	3053 CA ILE B	7	46.361		66.250	1.00 23.83
ATOM	3054 CB ILE B	7	45.736		66.670	
ATOM	3055 CG2 ILE B	7	44.309			1.00 25.11
ATOM	3056 CG1 ILE B	7	45.690		67.127	1.00 23.59
ATOM	3057 CD1 ILE B	ŕ			65.477	1.00 31.00
		7	47.021		64.906	1.00 38.60
ATOM			46.179		67.363	1.00 26.49
MOTA	3059 O ILE B	7	46.766	6.005	68.430	1.00 26.68

ATO	M 3060 N GLYB 8	45.	777 7 151		
ATON					1.00 29.5
ATON					1.00 30.2
			217 9.273	67.667	1.00 28.79
ATON			529 9.207	66.590	1.00 19.70
ATOM		44.0			1.00 26.4
ATOM	1- 3065 CA THR B 9				1.00 20.40
ATOM					1.00 29.3
ATOM					1.00 32.64
ATOM	. 7040	40.9			1.00 30.35
-	1 3068 CG2 THR B 9	41.4			1.00 33.42
ATOM	2000	43.4		69.302	1.00 33.42
ATOM		43.8		70.416	1.00 30.46
ATOM	3071 N LEU B 10	43.2	28 13.754		
ATOM		43.3			1.00 32.05
ATOM					1.00 34.75
ATOM		43.3		69.190	1.00 38.02
ATOM	2000	44.6		68.355	1.00 40.82
		44.9	61 15.472	67.394	1.00 43.62
MOTA		44.3	14 17.869	67.605	1.00 34.99
ATOM	3077 C LEU B 10	42.2		70.945	1.00 34.25
ATOM	3078 O LEU B 10	42.4			1.00 34.23
ATOM	3079 N ASP B 11	41.1		72.042	1.00 33.13
ATOM	3080 CA ASP B 11				1.00 30.61
ATOM	200	40.0		71.501	1.00 33.08
	2000	38.92		70.945	1.00 37.57
ATOM	3082 CG ASP B 11	38.37	12 13.643	69.621	1.00 43.14
ATOM	3083 OD1 ASP B 11	38.03	3 14.834	69.525	1.00 42.22
ATOM	3084 OD2 ASP B 11	38.28		68.681	1 00 45 50
ATOM	3085 C ASP B 11	40.35			1.00 45.58
MOTA	3086 O ASP B 11	39.68		72.919	1.00 32.19
ATOM	2000			73.875	1.00 23.44
ATOM	3000	41.38		73.06 6	1.00 28.02
	3000	41.77		74.402	1.00 32.00
ATOM	3089 CB TYR B 12	43.01	1 11.476	74.363	1.00 28.67
MOTA	3090 CG TYR B 12	42.82	1 10.108	73.737	1.00 25.33
MOTA	3091 CD1 TYR B 12	43.33		72.475	1.00 23.74
MOTA	3092 CE1 TYR B 12	43.23			1.00 23.74
ATOM	3093 CD2 TYR B 12	42.18		71.924	1.00 22.85
ATOM	3094 CE2 TYR B 12			74.436	1.00 21.93
ATOM	2005	42.07		73.889	1.00 21.99
ATOM	2004	42.60		72.640	1.00 22.99
	2000	42.53		72.109	1.00 18.79
ATOM	3097 C TYR B 12	42.05	4 13.567	75.319	1.00 32.74
ATOM	3098 O TYR B 12	41.98	6 13.450		1.00 23.85
ATOM	3099 N GLY B 13	42.37		74.720	1.00 26.96
ATOM	3100 CA GLY B 13	42.65		75.501	1 00 74 00
ATOM	3101 C GLY B 13	41.45		75.501	1.00 34.92
ATOM	3102 O GLY B 13				1.00 36.82
ATOM	7100	41.580			1.00 34.10
	2104	40.279		75.929	1.00 37.23
ATOM	22.00	39.031	. 16.247	76.584	1.00 41.77
ATOM	3105 CB LYS B 14	37.925	16.406	75.537	1.00 45.82
MOTA	3106 CG LYS B 14	38.110	17.585		1.00 51.38
ATOM	3107 CD LYS B 14	37.805	18.939		1.00 57.78
ATOM	3108 CE LYS B 14	38.752			
ATOM	3109 NZ LYS B 14	38.387	20 560		1.00 58.82
ATOM	3110 C LYS B 14	20 000			1.00 55.06
	2222 2 222 2 14	38.591		77.627	1.00 39.50
ATOM	2440	37.546	15.385	78.252 1	1.00 35.54
	3112 N TYR B 15	39.395	14.186		L.00 40.97
ATOM	3113 CA TYR B 15	39.070			.00 44.15
ATOM	3114 CB TYR B 15	38.863			
ATCM	3115 CG TYR B 15	37.850			.00 44.42
ATOM	3116 CD1 TYR B 15	30 054			.00 42.02
ATOM		38.064		75.634 1	.00 41.06
		37.138		74.603 1	.00 42.76
ATOM	3118 CD2 TYR B 15	36.678	12.703	77.065 1	.00 42.99
ATOM	3119 CE2 TYR B 15	35.748	12.851		.00 43.30
ATOM	3120 CZ TYR B 15	35.984			.00 45.49
ATOM	3121 OH TYR B 15	35.066		73.801 1	.00 45.69
ATOM	3122 C TYR B 15	40.151			00 43.69
ATOM	3123 O TYR B 15				.00 43.48
	3124 N ARG B 16	40.519			.00 41.20
ATCM		40.647		30.381 1	.00 43.01
atom	3125 CA ARG B 16	41.686	14.012	31.410 1	.00 43.70

MOTA	1 312	26 (CB ARG	B 16	5	42.25	0 15.41	0 81.66	3 1 00 40
ATOM	1 312	27 (CG ARG			42.65			
ATOM	1 312	28 (D ARG			43.85	8 15.62		
ATOM	312	1 6	VE ARG			44.30			
ATOM	313	0 0	Z ARG			44.62		-	
ATOM	313	1 N	WH1 ARG			44.55			
ATOM			TH2 ARG			45.02			
ATOM	. 313					41.09	_		
ATOM	313					39.88			
ATOM						41.94			
MOTA			A TYR			41.49			
MOTA	313		B TYR			42.500	. •		
ATOM	313		G TYR			42.413			
ATOM	313		D1 TYR			42.530			
ATOM	314	0 C	E1 TYR			42.502			
ATOM	314		D2 TYR			42.258			
ATOM	314:		E2 TYR			42.229			
ATOM	314:					42.355		84.210	
MOTA	3144					42.371			
MOTA	3145	5 C	TYR			41.377	–		
ATOM	3146	5 0	TYR			41.947			
ATOM	3147	7 N	PRO			40.647			
ATOM	3148	CI				39,958			
MOTA	3149) C2	PRO			40.448			1.00 45.33
ATOM	3150) CE	PRO	B 18		39.648			
MOTA	3151	. CG	PRO	B 18		40.096	13.015		
MOTA	3152	C	PRO	B 18		41.702	15.809		
MOTA	3153		PRO	B 18		42.789	15.244	88.317	
ATOM	3154		LYS			41.506	17.095	88.507	1.00 48.42
ATOM	3155					42.535	18.040	88.952	1.00 51.03
ATOM	3156					41.873	19.122	89.814	1.00 56.35
ATOM	3157	CC		_		40.630	18.657	90.563	1.00 65.69
ATOM	3158	CD				40.894	17.441	91.423	1.00 68.96
ATOM	3159	CE				39.602	16.882	91.999	1.00 71.85
MOTA	3160	NZ				39.825	15.603	92.731	1.00 72.79
ATOM ATOM	3161 3162	C	LYS I			43.830	17.593	89.639	1.00 48.62
ATOM	3163	NO	LYS I			44.912	18.009	89.235	1.00 49.04
ATOM	3164	CA	ASN E			43.745	16.775	90.678	1.00 43.99
ATOM	3165	CB	ASN E			44.957	16.356	91.375	1.00 43.86
ATOM	3166	CG	ASN E			44.740	16.440	92.890	1.00 45.92
ATOM	3167	OD:				44.418	17.848	93.355	1.00 49.44
ATOM	3168	ND2				45.194 43.268	18.779	93.138	1.00 47.72
ATOM	3169	c	ASN B			45.460	18.011	93.999	1.00 49.42
ATOM	3170	Õ	ASN B			46.496	14.960 14.521	91.008 91.494	1.00 39.49
ATOM	3171	N	HIS B			44.729	14.274	90.140	1.00 38.24
ATOM	3172	CA	HIS B			45.091	12.923	8 723	1.00 36.93 1.00 33.57
ATOM	3173	CB	HIS B			43.948	12.299	82.924	1.00 28.67
ATOM	3174	CG	HIS B	21		44.068	10.817	88.750	1.00 32.14
ATCM	3175	CD2	HIS B	21		44.779	10.076	87.867	1.00 26.15
ATOM	3176	ND1	HIS B	21			9.917		1.00 29.59
ATOM	3177	CE1	HIS B	21		43.743	8.686	89.212	1.00 23.65
atom	3178	NE2	HIS B	21		44.560	8.755	88.177	1.00 29.71
atom	3179	C	HIS B	21		46.348	12.928	88.852	1.00 29.06
MOTA	3180	0	HIS B	21		46.536	13.805	88.015	1.00 24.86
ATCM	3181	N	PRO B	22		47.225	11.937	89.035	1.00 30.50
ATOM	3182	CD	PRO B	22		47.187	10.802	89.976	1.00 31.51
ATOM	3183	CA	PRO B	22		48.446	11.880	88.231	1.00 29.58
ATOM	3184	CB	PRO B	22		49.055	10.549	88.656	1.00 33.72
ATOM	3185	CG	PRO B	22		43.658	10.489	90.124	1.00 31.07
TOM	3186	C	PRO B	22		48.176	11.950	86.728	1.00.28.52
MOT	3187	0	PRO B	22		48.989	12.474	85.972	1.00 31.85
TOM	3188	N	LEU B	23		47.030	11.435	86.297	1.00 24.47
TOM	3189	CA	LEU B	23		46.685	11.434	84.874	1.00 27.80
ATOM	3190	CB	LEU B	23		45.933	10.141	84.513	1.00 22.18
HTCH	3191	CG	LEU B	23		46.760	8.852	84.556	1.00 29.32

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ATOM	3192	e cr	1 LEU	J B	23	45.868	7.628	84.402	1.00 24.94
ATOM	3193				23	47.805		83.446	
ATOM	3194		LEU		23	45.891			
ATOM	3195		LEU		23			84.367	
ATOM	3196		LYS			45.166		83.373	1.00 24.42
					24	46.011		85.018	1.00 31.01
MOTA	3197				24	45.261		84.530	1.00 29.40
ATOM	3198				24	44.934		85.665	1.00 33.03
ATOM	3199				24	45.979	16.969	85.999	1.00 33.54
ATOM	3200				24	47.300	16.397	86.422	1.00 39.10
MOTA	3201		LYS	В	24	48.109	17.466	87.152	1.00 45.49
ATOM	3202	NZ	LYS	В	24	48.224	18.737	86.380	1.00 45.95
MOTA	3203	С	LYS	В	24	46.039	15.653	83.425	1.00 30.02
ATOM	3204	0	LYS	В	24	45.508	16.523	82.736	1.00 28.82
ATOM	3205	N	ILE	В	25	47.298	15.262	83.246	1.00 25.93
ATOM	3206	CA	ILE	В	25	48.139	15.858	82.212	1.00 29.48
ATOM	3207	CB	ILE		25	49.641	15.528	82.409	1.00 33.61
ATOM	3208	CG			25	50.126	16.033	83.775	1.00 32.27
ATOM	3209	CG:			25	49.851	14.014	82.263	
ATOM	3210				25	51.310	13.584		
ATOM	3211	. c	ILE		25			82.188	1.00 36.32
ATOM	3212	ō	ILE		25	47.784	15.318	80.834	1.00 30.08
	3213	N	PRO			47.263	14.210	80.704	1.00 25.37
MOTA					26	48.064	16.101	79.783	1.00 29.19
ATOM	3214	CD	PRO		26	48.650	17.448	79.770	1.00 32.47
ATOM	3215	CA	PRO		26	47.782	15.673	78.413	1.00 29.52
MOTA	3216	CB	PRO		26	48.103	16.921	77.593	1.00 29.84
MOTA	3217	CG	PRO		26	47.930	18.046	78.599	1.00 36.40
ATOM	3218	C	PRO		26	48.789	14.561	78.137	1.00 27.64
ATOM	3219	0	PRO		26	49.920	14.620	78.629	1.00 23.08
MOTA	3220	М	ARG		27	48.403	13.55 7	77 .360	1.00 23.09
ATOM	3221	CA	ARG	В	27	49.326	12 .469	77.072	1.00 23.00
ATOM	3222	CB		В	27	48.987	11.264	77.962	1.00 26.21
MOTA	3223	CG	ARG	В	27	49.101	11.617	79.449	1.00 17.03
MOTA	3224	CD	ARG	В	27	48.663	10.507	80.416	1.00 26.83
ATOM	3225	NE	ARG	В	27	49.586	9.375	80.502	1.00 22.99
ATOM	3226	CZ	ARG	В	27	49.444	8.220	79.856	1.00 25.06
ATOM	3227	NH1	ARG	В	27	48.408	8.022	79.059	1.00 17.74
ATOM	3228	NH2	ARG	В	27	50.336	7.253	80.027	1.00 23.38
MOTA	3229	C	ARG	В	27	49.329	12.097	7 5. 595	1.00 22.54
ATOM	3230	o	ARG	В	27	50.214	12.526	74.852	1.00 21.86
ATOM	3231	N	VAL	В	28	48.352	11.318	75.148	1.00 20.64
ATOM	3232	CA	VAL	B	28	48.337	10.954	73.739	1.00 26.57
ATOM	3233	CB	VAL	В	28	47.242	9.917	73.424	1.00 30.92
ATOM	3234	CG1	VAL	В	28	47.195	9.645	71.925	1.00 27.04
ATOM	3235	CG2	VAL		28	47.535	8.616	74.172	1.00 25.45
ATOM	3236	2	VAL		28	48.150	12.189	72.866	1.00 28.02
ATOM	3:37	Ö	VAL		28	48.780	12.311	71.808	1.00 30.88
ATOM	3.38	N	SER		29	47.298	13.112	73.304	1.00 24.30
ATOM	3139	CA	SER		29	47.082	14.326	72.523	1.00 29.48
ATOM	3240	CB	SER		29	45.939	15.169	73.110	1.00 29.48
ATOM	3241	OG	SER		29	46.218	15.614	74.424	
	3242	c	SER		29				1.00 34.55
ATOM	3243	5				48.379	15.125	72.514	1.00 30.81
ATOM	3244	27	SER :		29	48.680	15.820	71.545	1.00 28.85
ATOM	3245	CA			30	49.157	15.003	73.589	1.00 29.63
ATOM			LEU !		30	50.427	15.721	73.679	1.00 31.59
ATOM	3245	CB	LEU		30	51.046	15.593	75.079	1.00 29.49
ATOM	3247	CG	LEU		30	52.066	16.660	75.513	1.00 34.37
MOTA	3248		LEU 1		30	52.937	16.083	76.610	1.00 30.15
ATOM	3249		LEU I		30	52.951	17.098	74.357	1.00 32.90
ATOM	3250	2	LEU 1		30	51.371	15.085	7 2. 672	1.00-25.90
ATOM	3251	2	LEU I		30	52.052	15.777	71.913	1.00 25.10
ATOM	3252	N	LEU i		31	51.404	13.756	72.675	1.00 22.10
ATOM	3253	CA	LEU I		31	52.268	13.013	71.764	1.00 25.52
ATOM	3254	CB	LEU I		31	51.966	11.514	71.842	1.00 26.41
ATOM	3255	CG	LEU E	3	31	53.066	10.524	71.441	1.00 28.93
ATOM	3256	CD1	LEU E	3	31	52.425	9.198	71.042	1.00 23.69
ATOM	3257	CD2	LEU E	3	31	53.873	11.049	70.300	1.00 30.41

MOTA	3258	3 C	LEU	3 31	52.010	13.489	70.335	1.00 25.38
MOTA	3259	0	LEU		52.940		69.614	1.00 21.03
ATOM	3260	N	LEU		50.741		69.933	1.00 21.27
ATOM	3261	. CA	LEU		50.364		68.585	1.00 27.91
ATOM	3262	CB			48.841		68.408	1.00 26.60
ATOM	3263	CG			48.195		68.614	1.00 27.30
ATOM	3264		1-LEU		46.699		68.321	1.00 31.60
MOTA	3265		2 LEU		48.837		67.708	1.00 26.90
ATOM	3266			3 . 32	50.835		68.242	1.00 26.07
ATOM	3267		LEU		51.458		67.205	1.00 20.07
ATOM	3268		ARG		50.545		69.111	1.00 28.19
ATOM	. 3269				50.962		68.865	1.00 28.19
ATOM	3270		ARG		50.395		69.930	1.00 34.22
ATOM	3271				48.887	18.740	69.904	1.00 40.33
ATOM	3272	CD			48.420		70.970	1.00 47.67
ATOM	3273	NE		B 33	46.977	19.931	70.924	1.00 56.24
ATOM	3274	CZ	ARG		46.330	20.505	69.912	1.00 60.10
ATOM	3275	NH			46.997	20.929	68.845	1.00 63.11
MOTA	3276	NH			45.011	20.652	69.965	1.00 63.81
ATOM	3277	C	ARG		52.476	17.791	68.852	1.00 30.12
ATOM	3278	0	ARG		53.028	18.580	68.097	1.00 30.20
ATOM	3279	N	PHE		53.147	17.012	69.694	1.00 30.70
ATOM	3280	CA	PHE		54.600	17.060	69.774	1.00 29.42
ATOM	3281	CB	PHE		55.096	16.176	70.920	1.00 30.46
ATOM	3282	CG	PHE		56.556	16.358	71.248	1.00 28.56
ATOM	3283	CDI	PHE		57.001	17.515	71.885	1.00 26.92
MOTA	3284	CD2	PHE	3 3 4	57.481	15.373	70.932	1.00 28.88
MOTA	3285	CE1	PHE	B 34	58.346	17.684	72.206	1.00 28.15
ATOM	3286	CE2	PHE	B 34	58.831	15.530	71.246	1.00 31.47
MOTA	3287	CZ	PHE	B 34	59.265	16.689	71.887	1.00 28.15
MOTA	3288	С	PHE .	B 34	55.202	16.583	68,460	1.00 33.78
MOTA	3289	0	PHE	B 34	56.049	17.259	67.873	1.00 33.71
ATOM	3290	N	LYS		54.770	15.413	67.999	1.00 28.65
ATOM	3291	CA	LYS		55.294	14.880	66.753	1.00 34.33
ATOM	3292	СВ	LYS :		54.684	13.509	66.454	1.00 32.97
MOTA	3293	CG	LYS		55.141	12.423	67.414	1.00 34.93
ATOM	3294	CD	LYS		54.580	11.066	67.047	1.00 41.43
MOTA	3295	CE	LYS		53.070	11.004	67.205	1.00 44.04
ATOM	3296	NZ	LYS		52.335	11.984	66.345	1.00 60.09
ATOM	3297	C	LYS !		55.015	15.842	65.608	1.00 35.78
MOTA	3298	0	LYS !		55.869	16.061	64.752	1.00 33.39
ATOM	3299	N	ASP 1		53.823	16.426	65.602	1.00 32.32
ATOM	3300 3301	CA CB	ASP S		53.468	17.365	64.552	1.00 36.31
ATOM	3301	CG	ASP E		52.015	17.800	64.698	1.00 42.56
ATOM	3302	OD1			51.617 51.812	18.822	63.661	1.00 43.03
MOTA	3304	OD2	ASP E			18.544	62.461 64.043	1.00 79.17
ATOM ATOM	3305	C	ASP E		51.111 54.371	19.897		1.00 .4.34
ATOM	3306	ō	ASP E		54.764	18.590 19.099	64.578 63.534	1.00 '6.14 1.00 32.40
ATOM	3307	N	ALA E		54.694	19.061	65.777	1.00 34.80
ATOM	3308	CA	ALA E		55.554	20.226		1.00 36.82
ATOM	3309	CB	ALA B		55.599	20.220	67.383	1.00 38.54
MOTA	3310	c	ALA B		56.959	19.901	65.429	1.00 37.66
ATOM	3311	ō	ALA B		57.675	20.776	64.950	1.00 30.56
ATOM	3312	N	MET B		57.346	18.635	65.541	1.00 37.42
MOTA	3313	CA	MET 3		58.670	18.192	65.107	1.00 37.42
ATOM	3314	CB	MET B		59.158	17.059	66.013	1.00 36.44
ATOM	3315	CG	MET 3		59.341	17.438	67.474	1.00 37.68
MOTA	3316	SD.	MET 3		60.841	18.391	67.784	1.00 38.07
ATOM	3317	CE .	MET 3	_	62.093	17.228	67.300	1.00 30.98
ATOM	3318	c_	MET B		58.639	17.690	63.663	1.00 35.86
ATOM	3319	ō	MET B		59.659	17.262	63.130	1.00 32.69
ATOM	3320	N	ASN B		57.470	17.742	63.035	1.00 35.82
ATOM	3321	CA	ASN B	39	57.321	17.262	61.661	1.00 42.75
ATOM	3322	CB	ASN B	39	58.156	18.108	60.688	1.00 46.20
ATOM	3323	CG	ASN B	39	57.670	19.543	60.591	1.00 47.57
					•			

ATOM	1 332	4 0	D1 AS	N I	3 39	56.52	4 19.80	1 60.21	2 1 00 40 70
ATOM	332		D2 AS			58.54			
ATOM	332			N E		57.75			
ATOM	332			N E		58.46			
ATOM	332	8 N		UE		57.33			
ATOM	332	9 C2		UE		57.70			
ATOM				UE		58.34			
ATOM				UB		59.59			
MOTA			01 LE			60.14			
ATOM			02 - LEI			60.64			
ATOM			LE			56.549			
ATOM			LE			56.637			
ATOM				В		55.476			
ATOM	3337			5 B		54.340			
ATOM	3338				41	53.445			
ATOM	3339				41	52.793			
ATOM	3340				41	52.367			
ATOM	3341	. CD			41	51.470			
MOTA	3342	С	ILE		41	53.492			
ATOM	3343	0	ILE		41	53.352			
ATOM	3344	N	ASP		42	52.943			
ATOM	3345				42	52.094	12.615		
ATOM	3346	CB			42	52.569	12.119		
ATOM	3347	CG	ASP	В	42	53.972	12.584		
ATOM	3348	OD:	1 ASP	В	42	54.244	13.799		
MOTA	3349	OD:	2 ASP	В	42	54.797	11.736		1.00 45.16
MOTA	3350	С	ÄSP	В	42	50.677	12.134		1.00 45.15
ATOM	3351	0	ASP	В	42	50.467	11.051		1.00 47.06
ATCM	3352	N	GLU	В	43	49.707	12.944		1.00 48.13
MOTA	3353	CA	GLU	В	43	48.303	12.618		1.00 50.50
ATOM	3354	CB	GLU		43	47.441	13.637	57.571	1.00 53.54
ATOM	3355	CG	GLU		43	45.961	13.505	57.840	1.00 59.52
MOTA	3356	CD	GLU		43	45.155	14.518	57.065	1.00 64.03
ATOM	3357	OE1			43 .	43.914	14.535	57.215	1.00 68.54
ATOM	3358	OE2			43	45.765	15.298	56.301	1.00 66.95
MOTA	3359	C	GLU		43	47.972	11.205	57.836	1.00 47.81
ATOM	3360	0	GLU		43	47.092	10.547	58.390	1.00 49.67
ATOM	3361 3362	N	LYS		44	48.690	10.744	56.817	1.00 46.21
ATOM ATOM	3362	CA	LYS		44	48.484	9.409	56.251	1.00 48.28
ATOM	3364	CB	LYS LYS		44	49.207	9.311	54.894	1.00 49.96
ATOM	3365	CD	LYS		44	49.639	7.903	54.470	1.00 52.18
ATOM	3366	CE	LYS		44 44	50.970	7.532	55.127	1.00 61.03
ATOM	3367	NZ	LYS	-	44	51.399	6.095	54.844	1.00 62.80
ATOM	3368	C	LYS		44	50.511	5.098	55.510	1.00 65.34
ATOM	3369	ō	LYS		44	48.899	8.249	57.161	1.00 45.92
MOTA	3370	N	GLU		45	48.418 49.797	7.127	57.009	1.00 41.30
ATOM	3371	CA	GLU		45	50.268	8.517 7.486	581100 59.014	1.00 42.18
ATOM	3372	CB	GLU		45	51.684	7.812	59.468	1.00 38.41 1.00 33.73
ATOM	3373	CG	GLU :		45	52.694	7.812	58.351	
ATOM	3374	CD	GLU :		45	53.998	8.504	58.813	1.00 37.58
MOTA	3375		GLU		45	53.997	9.699	59.176	1.00 34.34 1.00 38.04
ATOM	3376		GLU I		45	55.020	7.799	58.821	1.00 33.37
ATOM	3377	C	GLU I		45	49.368	7.403	60.238	1.00 36.86
ATOM	3378	0	GLU I		45	49.461	6.462	61.032	1.00 34.98
ATCM	3379	N	LEU I		46	48.489	8.386	60.386	1.00 30.86
ATOM	3380	CA	LEU H	3	46	47.608	8.438	61.545	1.00 30.65
MOTA	3381		LEU E		46	47.501	9.889	62.019	1.00 30.03
MOTA	3382		LEU I		46	46.642	10.163	63.250	1.00 32.74
ATOM	3383		LEU F		46	47.189	9.379	64.425	1.00 34.70
ATOM	3384		LEU E		46	46.639	11.656	63.548	1.00 33.94
ATOM	3385	С	LEU F		46	46.212	7.861	61.318	1.00 31.36
MOTA	3386		LEU F		46	45.530	8.218	60.363	1.00 31.78
ATOM	3387		ILE E		47	45.801	6.957	62.203	1.00 31.18
ATOM	3388		ILE E		47	44.479	6.338	62.139	1.00 29.36
ATOM	3389	CB	ILE E	3 .	47	44.564	4.802	62.258	1.00 28.62

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Figure 18-52

MOTA MOTA MOTA MOTA MOTA MOTA MOTA MOTA	339 339 339 339 339 339 339 3400 3400 34	1 CG1 ILE 2 CD1 ILE 3 C ILE 4 O ILE 5 N LYS 6 CA LYS 7 CB LYS 7 CB LYS 8 CD LYS 9 CD LYS 10 CE LYS 11 NZ LYS 12 C LYS 13 C LYS 15 C LYS 16 C LYS 17 CE LYS 18 C LYS 1	B 47 B 47 B 47 B 47 B 48 B 48 B 48 B 48 B 48 B 48 B 48		43.163 45.266 45.419 43.659 44.063 42.514 41.662 40.517 39.607 38.535 37.657 38.451 41.095	4.230 2.722 6.875 6.755 4.7.475 8.840 9.514 6.10.361 11.074 11.991	61.028 61.054 63.303 64.461 62.999	1.00 29.42 1.00 31.12 1.00 32.22 1.00 31.17 1.00 28.39 1.00 32.37 1.00 36.32 1.00 44.38 1.00 45.91 1.00 42.66
MOTA	3403 3404		-		40.524 41.260	5.962	64.457 66.244	1.00 26.24
ATOM	3405	CA SER'	-,-		40.770		67.232	1.00 27.89 1.00 25.17
MOTA	3406				41.146		68.642	1.00 24.41
ATOM ATOM	3407 3408				42.539 39.248		68.777	1.00 31.79
ATOM	3409				38.565		67.160 66.879	1.00 29.07 1.00 28.47
MOTA	3410				38.723	4.859	67.409	1.00 26.13
ATOM ATOM	3411 3412	CA ARG E			37.278	4.658	67.430	1.00 24.24
ATOM	3413	CG ARG E			36.810 37.231	3.700 2.233	66.323 66.507	1.00 25.03 1.00 26.54
ATOM	3414	CD ARG B	50		36.570	1.340	65.452	1.00 26.21
ATOM ATOM	3415 3416	NE ARG B			37.006	-0.058	65.504	1.00 25.13
ATOM	3417	NH1 ARG B			36.700 35.941	-0.924 -0.558	66.468 67.497	1.00 26.09 1.00 23.42
ATOM	3418	NH2 ARG B	50		37.157	-2.168	66.402	1.00 23.91
ATOM ATOM	3419 3420	C ARG E O ARG B			36.937	4.037	68.775	1.00 23.83
ATOM	3421	N PRO B	50 51		37.782	3.392 4.223	69.403 69.243	1.00 21.60
ATOM	3422	CD PRO B	51		34.554	4.962.		1.00 25.09
ATOM ATOM	3423 3424	CA PRO B	51		35.338	3.628	70.530	1.00 25.48
ATOM	3425	CG PRO B	51 51		33.949 33.936	4.217 5.503	70.802 69.953	1.00 26.32 1.00 28.65
ATOM	3426	C PRO B	51		35.264	2.118	70.325	1.00 26.73
ATOM ATOM	3427 3428	O PROB	51 52		35.142	1.646	69.194	1.00 18.87
ATOM	3429	CA ALA B	52		35.355 35.237	1.359 -0.083	71.408 71.291	1.00 23.64 1.00 23.27
ATOM	3430	CB ALA B	52		35.811	-0.757	72.521	1.00 26.31
atom atom	3431 3432	C ALA B	52 52		33.733	-0.324	71.223	1.00 25.25
ATOM	3433	N THR B	53		32.950 33.321	0.515 -1.447	71.677 70.651	1.00 22.78 1.00 22.77
ATOM	3434	CA THR B	53 ·		31.900	-1.760	70.596	1.00 26.90
MOTA MOTA	3435 3436	CB THR B	53 53		31.567	-2.732	69.456	1.00 30.00
ATOM	3437	CG2 THR B	53	•	32.305 31.917	-3.950 -2.117	69.642 68.103	1.00 25.59 1.00 23.33
MOTA	3438	C THR B	53		31.579	-2.445	71.916	1.00 30.41
MOTA MOTA	3439 3440	O THR B N LYS B	53 54		32.484	-2.917	72.609	1.00 26.13
ATOM	3441	CA LYS B	54		30.300 29.909	-2.504 -3.140	72.268 73.514	1.00 29.12 .
ATOM	3442	CB LYS B	54		28.396	-3.027	73.720	1.00 32.78
ATOM	3443 3444	CG LYS B	54		27.947	-3.351	75.131	1.00 34.85
ATOM ATOM	3445	CD LYS B	54 54		26.445 26.008	-3.204 -3.366	75.268 76.709	1.00 41.13 1.00 43.39
ATOM	3446	NZ LYS B	54		26.464	-2.257	77.582	1.00 47.39
ATOM	3447	C LYS B	54		30.329	-4.603	73.442	1.00.29.10
atom atom	3448 3449	O LYS B	54 55		30.779 30.196	-5.183 -5.187	74.430 72.256	1.00 26.71 1.00 23.97
ATOM	3450	CA GLU B	55		30.136	-5.107 -6.577	72.236	1.00 28.08
ATCM	3451	CB GLU B	55		30.288	-6.965	70.579	1:00 24.82
ATOM ATOM	3452 3453	CG GLU B	55 55		30.671 30.453	-8.400	70.237 68.767	1.00 33.40 1.00 38.49
ATOM	3454	OE1 GLU B	55		30.638	-8.737 -9.913	68.394	1.00 38.49
ATOM	3455	OE2 GLU B	55		30.101	-7.833	67.984	1.00 40.02

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ATOM	3456	5 C	GLU	B 55		32.066	-6.808	72.345	1 00 25 02
	3457								1.00 25.82
ATOM			GLU			32.429		73.033	1.00 23.83
ATOM	3458	3 N	GLU	B 56		32.931	5.938	71.835	1.00 25.04
ATOM	3459) CF	A GLU			34.365		72.093	
									1.00 25.30
ATOM	3460					35.141	-5.003	71.334	1.00 24.31
ATOM-	3461	. CG	GLU	B 56		34.866	-5.039	69.836	1.00 32.15
ATOM	3462	CI				35.512			
								69.073	1.00 31.43
MOTA	3463	OE	:1 GLU	. В 56		35.486	-2.759	69.568	1.00 28.54
MOTA	3464	OE	:2 GLU	B 56		36.012	-4.147	67.959	1.00 28.89
ATOM	3465		GLU						
						34.653		73.595	1.00 28.88
ATOM	3466	0	GLU	B 56		35.450	-6.766	74.137	1.00 25.07
ATOM	3467	N	LEU	B. 57		33.996	-5.050	74.272	1.00 24.52
ATOM	3468				-				
						34.203		75.702-	1.00 27.34
MOTA	3469	CB	LEU	B 57		33.416	-3.694	76.231	1.00 22.79
ATOM	3470	CG	LEU	B 57		33.859	-2.320	75.722	1.00 23.57
ATOM	3471	CD							
						33.008		76.366	1.00 22.27
MOTA	3472	CD	2 LEU	B 57		35.342	-2.089	76.061	1.00 17.24
MOTA	3473	С	LEU	B 57		33.785	-6.144	76.452	1.00 26.92
ATOM	3474	0	LEU			34.458			
								77.396	1.00 24.06
MOTA	3475	N	LEU	B 58		32.670	-6.732	76.029	1.00 23.35
MOTA	3476	CA	LEU	B 58		32.154	-7.931	76.674	1.00 25.60
ATOM	3477	CB	LEU			30.718			
							-8.207	76.221	1.00 28.50
ATOM	3478	CG	LEU	B 58		29.734	-7.110	76.649	1.00 30.91
ATOM	3479	CD:	1 LEU	B 58		28.323	-7.468	76.212	1.00 28.93
ATOM	3480	CD:	2 LEU			29.794	-6.945	78.157	1.00 33.44
ATOM	3481	С	LEU			33.027	- 9.153	76.446	1.00 24.59
ATOM	3482	0	LEU	B 58		32.760	-10.216	76.991	1.00 19.76
ATOM	3483	N	LEU	B 59		34.065	-9.006	75.630	1.00 23.99
	3484	CA	LEU						
MOTA							-10.108	75.411	1.00 25.11
ATOM	3485	CB	LEU	B 59		36.018	-9.757	74.332	1.00 21.64
ATOM	3486	CG	LEU	B 59		35.483	-9.652	72.905	1.00 24.24
ATOM	3487	CDI				36.585			
							-9.177	71.975	1.00 24.25
MOTA	3488	CD2	LEU	B 59		34.957	-11.014	72.468	1.00 19.91
ATOM	3489	С	LEU :	B 59		35.699	-10.371	76.733	1.00 23.84
MOTA	3490	0	LEU :				-11.489	76.992	1.00 19.39
atom	3491	N	PHE 1			35.793	-9.344	77.577	1.00 21.80
ATOM	3492	CA	PHE 1	в 60		36.462	-9.510	78.876	1.00 23.08
ATOM	3493	CB	PHE I	B 60		37.809	-8.770	78.908	1.00 18.22
	3494	CG	PHE						
ATOM						38.544	-8.906	80.230	1.00 21.72
ATOM	3495	CD1	PHE 1	в 60		38.975	-10.157	80.680	1.00 19.23
ATOM	3496	CD2	PHE I	B 60		38.757	-7.791	81.048	1.00 17.75
ATOM	3497	CE1					-10.301	81.927	1.00 18.80
atom	3498	CE2				39.384	-7.923	82.297	1.00 19.23
ATOM	3499	CZ	PHE B	3 60 E		39.807	-9.184	82.737	1.00 16.10
ATOM	3500	С	PHE E			35.648	-9.069	80.083	1.00 21.58
ATOM	3501	0	PHE F			35.508	-9.822	81.040	1.00 22.21
ATOM	3502	N	HIS E	3 61		35.128	~7.847	80.055	1.00 20.65
ATOM	3503	CA	HIS E	61		34.362	-7.336	81.184	1.00 23.32
ATOM	3504	СВ	HIS E					81.229	
						34.422	-5.807		1.00 27.60
ATOM	3505	CG	HIS E			35.800	-5.259	81.440	1.00 31.83
ATOM	3506	CD2	HIS E	61		36.466	-4.940	82.575	1.00 26.86
	3507		HIS B				~5.003	80.401	
ATOM						36.669			1.00 34.35
ATOM	3508	CEI	HIS B	61		37.810	-4.546	80.887	1.00 34.78
ATOM	3509	NE2	HIS B	61		37.713	-4.499	82.204	1.00 36.27
ATOM	3510	С	HIS B			32.902	-7.775	81.198	1.00 28.04
atcm	3511	0	HIS B			32.349	-8.167	80.176	1.00 25.70
ATOM	3512	N	THR B	62		32.276	-7.691	82.367	1.00 25.25
ATOM	3513	CA	THR B			30.882	-8.084	82.506	1.00 25.35
									2 00 05 45
ATOM	3514	CB	THR B			30.578	-8.549	83.932	1.00 25.47
ATOM	3515	OG1	THR B	62		30.783	-7.462	84.843	1.00 28.62
ATOM	3516	CG2	THR B	62		31.482	-9.701	84.315	1.00 21.32
		c	THR B			29.931		82.162	
ATOM	3517						-6.942		1.00 26.06
ATOM	3518	0	THR B			30.287	-5.771	82.254	1.00 24.14
ATCM	3519	N	GLU B	63		28.718	-7.305	81.759	1.00 28.01
	3520	CA	GLU B			27.681		81.389	1.00 30.77
ATCM							-6.349		
ATCM	3521	CB	GLU B	63		26.374	-7.094	81.114	1.00 33.97

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				•			
MOTA	3522	CG .GLU	B 63	25 21	2 (22)		_
							7 1.00 41.12
ATOM		CD GLU		25.18	9 ~5.98	7 79.168	
ATOM	3524	OE1 GLU	B 63				
ATOM	3525	OE2 GLU					
	_		_				1.00 45.96
ATOM		C GLU		27.43	5 -5.326	82.498	
ATOM	3527	O GLU	B 63	27.383			
ATOM	3528	N ASP					
							1.00 24.38
ATOM		CA ASP	B .64	27.010	-5.023	84.897	1.00 29.27
ATOM	3530	CB ASP	B 64	26.887			
ATOM	3531	CG ASP					
				28.022		86.198	1.00 50.77
MOTA		OD1 ASP	B 64	29.128	-6.540	86.630	
MOTA	3533	OD2 ASP	B 64	27.812			
ATOM		C ASP					
				. 28.075		85.143	1.00 26.49
ATOM		O ASP	B 64	27.768	-2.806	85.422	1.00 18.33
ATOM	3536 1	Y TYR	B 65	29.332			
ATOM	3537	CA TYR					
			_	30.420			1.00 19.32
ATOM		CB TYR :	B 65	31.751	-4.186		1.00 16.59
ATOM	3539 (G TYR	B 65	32.949			
ATOM		D1 TYR	-				
				33.033		86.383	1.00 21.35
ATOM		CE1 TYR 1		34.135	-1.489	86.489	1.00 18.32
ATOM	3542	D2 TYR I	B 65	34.004	-3.382	84.456	1 00 10.32
ATOM		E2 TYR I					1.00 18.65
				35.116	-2.544	84.554	1.00 21.01
ATOM ·		Z TYR I		35.172	-1.601	85.573	1.00 20.61
ATOM	3545 C	H TYR E	3 65	36.262	-0.775		
ATOM	3546 C						1.00 17.77
				30.392	-2.373	84.146	1.00 22.01
ATOM				30.399	-1.167	84.421	1.00 18.20
ATOM	3548 N	ILE E	66	30.330	-2.815	82.894	1.00 19.49
ATOM	3549 C	A ILE E		. 30.305			
ATOM	3550 C				-1.870	81.786	1.00 19.68
				30.208	-2.592	80.432	1.00 23.31
ATOM		G2 ILE B		30.200	-1.571	79.303	1.00 21.30
ATOM	3552 C	G1 ILE B	66	31.400	-3.541		
ATOM		D1 ILE B				80.260	1:00 27.67
		. –		32.758	-2.839	80.291	1.00 29.29
ATOM	3554 C	ILE B	66	29.128	-0.909	81.940	1.00 26.99
ATOM	3555 o	ILE B		29.294	0.309	81.848	1.00 23.36
ATOM	3556 N	ASN B	67	27.939			
ATOM	3557 C				-1.447	82.198	1.00 24.98
			67	26.782	~0.580	82.363	1.00 27.70
ATOM	3558 CI	B ASN B	67	25.492	~1.389	82.580	1.00 25.58
MOTA	3559 CC	ASN B	67	25.081	-2.183	01 341	
MOTA		Ol ASN B	67			81.341	1.00 26.91
				25.199	~1.701	80.220	1.00 31.48
MOTA	3561 NI		67	24.572	-3.387	81.545	1.00 23.80
ATOM	3562 C	ASN B	67	26.982	0.401	83.513	1.00 25.34
ATOM	3563 O	ASN B	67	26.524			
ATOM	3564 N				1.539	83.448	1.00 22.53
		THR B	68	27.664	-0.031	84.568	1.00 23.65
ATOM	3565 CA		68	27.903	0.863	85.696	1.00 25.25
ATOM.	3566 CB	THR B	68	28.516	0.119		
ATOM .	3567 OG	1 THR B	68			86.891	1.00 29.08
				27.561	-0.826	87.396	1.00 25.94
MOTA	3568 CG		68	28.894	1.100	88.002	1.00 22.90
ATOM	3569 C	THR B	68	28.818	2.009	85.287	1.00 25.91
ATOM	3570 O	THR B	68	28.576	3.156	85.661	
ATOM	3571 N	LEU B					1.00 28.47
			69	29.861	1.702	84.519	1.00 25.13
ATOM	3572 CA	LEU B	6 <i>9</i>	30.788	2.729	84.054	1.00 24.37
MOTA	3573 CB	LEU B	69	31.915	2.122		
ATOM	3574 CG	LEU B	69			83.201	1.00 21.32
		1000		32.960	1.231	83.889	1.00 22.33
ATOM	3575 CD	LEU B	69	34.006	0.786	82.859	1.00 22.57
ATOM	3576 CD2	LEU B	69	33.643	2.000		
ATOM	3577 C	LEU B	69	30.036			1.00 23.20
					3.764	83.229	1.00 23.02
• • • • • • •	3578 O	LEU B	69	30.190	4.966	83.444	1.00 18.98
MOTA	3579 พ	MET B	70	29.218	3.290		1.00 19.62
	3580 CA	MET B	70	28.449			
					4.181	81.434	1.00 25.87
	3581 CB	MET B	70	27.660	3.371	80.401	1.00 24.80
ATOM	3582 CG	MET B	70	28.531	2.511		1.00 30.37
	3583 SD	MET B	70	27.592			
							1.00 30.35
		MET B	70	26.922	2.986	77.245	1.00 30.20
ATOM	3585 C	MET B	70	27.489		1	1.00 28.82
	3586 O	MET B	70	27.391			
							1.00 24.09
MOTA	3587 N	GLU B	71	26.786	4.458	83.194	1.00 28.21

ATOM 3588 CA GLU B 71										
ATOM 3589 CG GLU B 71	MOTA	358	B CA G	LU I	3 71		25.837	5.207	84.008	1.00 27.45
ATOM 3590 CG GLU B 71 24.072 5.005 85.832 1.00 33.61 ATOM 3591 CD GLU B 71 23.044 5.867 85.096 1.00 37.51 ATOM 3593 OEL GLU B 71 22.333 6.638 85.773 1.00 35.47 ATOM 3593 OEL GLU B 71 22.333 6.638 85.773 1.00 35.47 ATOM 3594 C GLU B 71 22.333 6.638 85.773 1.00 35.47 ATOM 3595 O GLU B 71 22.333 6.638 85.773 1.00 29.15 ATOM 3595 O GLU B 71 26.559 6.209 84.887 1.00 29.15 ATOM 3596 N ALA B 72 27.671 5.781 85.481 1.00 27.76 ATOM 3597 C ALA B 72 28.454 6.662 86.340 1.00 27.58 ATOM 3598 CB ALA B 72 28.454 6.662 86.340 1.00 27.58 ATOM 3599 C ALA B 72 28.895 8.999 86.079 1.00 23.24 ATOM 3600 O ALA B 72 28.895 8.999 86.079 1.00 23.24 ATOM 3601 N GLU B 73 29.356 7.684 84.322 1.00 29.48 ATOM 3602 CA GLU B 73 30.658 8.314 82.325 1.00 29.48 ATOM 3603 CB GLU B 73 31.162 9.466 81.443 1.00 31.03 ATOM 3604 CG GLU B 73 31.938 9.009 80.216 1.00 34.37 ATOM 3607 OEZ GLU B 73 31.938 9.009 80.216 1.00 34.37 ATOM 3608 C GLU B 73 31.938 9.009 80.216 1.00 34.37 ATOM 3607 OEZ GLU B 73 31.419 9.203 79.100 1.00 30.59 ATOM 3608 C GLU B 73 31.419 9.203 79.100 1.00 30.59 ATOM 3609 O GLU B 73 31.419 9.203 79.100 1.00 30.59 ATOM 3601 N ARG B 74 26.583 10.067 82.081 1.00 31.92 ATOM 3601 N ARG B 74 27.633 9.186 82.570 1.00 39.90 ATOM 3611 CA ARG B 74 22.548 8.91 83.294 1.00 53.58 ATOM 3612 CB ARG B 74 22.142 1.0351 83.297 1.00 29.48 ATOM 3613 CG ARG B 74 22.142 8.311 81.646 1.00 31.92 ATOM 3614 CD ARG B 74 22.142 8.311 81.646 1.00 35.84 ATOM 3615 NE ARG B 74 22.142 8.311 81.646 1.00 35.84 ATOM 3616 NE ARG B 74 22.142 8.311 81.646 1.00 35.84 ATOM 3617 NH1 ARG B 74 22.142 8.311 81.646 1.00 34.37 ATOM 3620 C ARG B 74 22.142 8.311 81.646 1.00 34.37 ATOM 3621 C B ARG B 74 22.142 8.311 81.646 1.00 39.90 ATOM 3622 C ARG B 74 22.142 8.311 81.646 1.00 39.90 ATOM 3631 C C GLU B 73 31.419 9.203 84.386 1.00 33.58 ATOM 3632 C G G G G G G G G G G G G G G G G G G		3589	CB G	LU I	3 71					
ATOM 3591 CD GLU B 71		3590	CG G	LU I	3 71					
ATOM 3592 OEL GLU B 71 22.333 6.638 85.773 1.00 35.47 ATOM 3593 OE2 GLU B 71 22.934 5.769 83.849 1.00 31.03 ATOM 3595 C GLU B 71 26.559 6.209 84.887 1.00 29.15 ATOM 3595 O GLU B 71 26.559 6.209 84.887 1.00 29.15 ATOM 3595 O GLU B 71 26.115 7.341 85.035 1.00 23.96 ATOM 3596 N ALA B 72 28.454 6.662 86.340 1.00 27.58 ATOM 3598 CB ALA B 72 28.454 6.662 86.340 1.00 27.58 ATOM 3599 CA ALA B 72 29.663 5.920 86.909 1.00 23.24 ATOM 3590 C ALA B 72 28.895 8.999 86.079 1.00 23.24 ATOM 3600 O ALA B 72 28.895 8.999 86.079 1.00 23.24 ATOM 3601 N GLU B 73 29.356 7.684 84.322 1.00 26.40 ATOM 3602 CA GLU B 73 29.356 7.684 84.322 1.00 29.48 ATOM 3603 CB GLU B 73 30.658 8.314 82.325 1.00 29.48 ATOM 3604 CG GLU B 73 31.938 9.009 80.216 1.00 34.37 ATOM 3605 CD GLU B 73 31.938 9.009 80.216 1.00 34.37 ATOM 3607 OE2 GLU B 73 31.938 9.009 80.216 1.00 34.37 ATOM 3607 OE2 GLU B 73 31.938 9.009 80.216 1.00 36.59 ATOM 3609 O GLU B 73 31.419 9.203 79.100 1.00 30.59 ATOM 3601 N ARG B 74 27.633 9.186 82.570 1.00 29.08 ATOM 3611 CA ARG B 74 26.583 10.067 82.081 1.00 38.64 ATOM 3612 CB ARG B 74 26.583 10.067 82.081 1.00 38.64 ATOM 3612 CB ARG B 74 22.448 8.706 82.365 1.00 46.67 ATOM 3612 CB ARG B 74 22.448 8.706 82.365 1.00 46.67 ATOM 3613 CG ARG B 74 22.448 8.706 82.365 1.00 46.67 ATOM 3615 NE ARG B 74 22.448 8.706 82.365 1.00 46.67 ATOM 3613 CG ARG B 74 22.448 8.706 82.365 1.00 46.67 ATOM 3613 CG ARG B 74 22.448 8.706 82.365 1.00 46.67 ATOM 3613 CG ARG B 74 22.448 8.706 82.365 1.00 55.58 ATOM 3618 NH2 ARG B 74 22.142 10.351 83.222 1.00 35.84 ATOM 3622 CB ARG B 74 22.448 8.706 82.365 1.00 36.99 ATOM 3622 CB ARG B 74 22.448 8.706 82.365 1.00 36.99 ATOM 3623 CB SER B 75 25.581 9.228 87.123 1.00 33.43 ATOM 3628 CB GRG B 74 22.468 8.706 82.365 1.00 35.84 ATOM 3628 CB GRG B 74 22.468 8.706 82.365 1.00 35.84 ATOM 3623 CB SER B 75 25.581 9.228 87.123 1.00 33.45 ATOM 3628 CB GRG B 74 22.468 8.706 82.365 1.00 35.84 ATOM 3628 CB GRG B 74 22.468 8.706 82.365 1.00 35.84 ATOM 3628 CB GRG B 75 25.581 9.228 87.123 1.00 33.45 ATOM 3628 CB GLB B 75 26.339 11.7		3591	L CD G	LU E	3 71					
ATOM 3593 OE2 GLU B 71 22.934 5.769 83.849 1.00 31.03 ATOM 3595 O GLU B 71 26.515 6.209 84.887 1.00 23.96 ATOM 3595 O GLU B 71 26.515 7.341 85.035 1.00 23.96 ATOM 3596 N ALA B. 72 27.671 5.781 85.481 1.00 27.76 ATOM 3597 CA ALA B 72 28.924 7.886 85.563 1.00 23.24 ATOM 3599 C ALA B 72 28.924 7.886 85.663 1.00 23.24 ATOM 3599 C ALA B 72 28.924 7.886 85.563 1.00 23.24 ATOM 3600 O ALA B 72 28.924 7.886 85.563 1.00 23.24 ATOM 3600 O ALA B 72 28.895 8.999 86.079 1.00 23.24 ATOM 3600 CA GLU B 73 29.846 8.801 83.529 1.00 29.06 ATOM 3603 CB GLU B 73 30.658 8.314 82.325 1.00 29.06 ATOM 3605 CD GLU B 73 31.938 9.009 80.216 1.00 31.00 ATOM 3605 CD GLU B 73 31.938 9.009 80.216 1.00 34.37 ATOM 3605 CD GLU B 73 31.938 9.009 80.216 1.00 28.41 ATOM 3608 C GLU B 73 31.938 9.009 80.216 1.00 28.41 ATOM 3608 C GLU B 73 31.419 9.203 79.100 1.00 35.69 ATOM 3601 N ARG B 74 27.633 9.186 82.570 1.00 28.41 ATOM 3601 C GLU B 73 31.419 9.203 79.100 1.00 35.59 ATOM 3601 C GLU B 73 31.419 9.203 79.100 1.00 35.59 ATOM 3601 C GLU B 73 28.844 10.951 83.104 1.00 31.92 ATOM 3608 C GLU B 73 28.744 9.734 83.045 1.00 28.41 ATOM 3608 C GLU B 73 28.744 9.734 83.045 1.00 31.92 ATOM 3601 C ARG B 74 27.633 9.186 82.570 1.00 35.59 ATOM 3601 C ARG B 74 27.633 9.186 82.570 1.00 35.59 ATOM 3611 C ARG B 74 27.633 9.186 82.570 1.00 35.53 ATOM 3611 C ARG B 74 26.583 10.067 82.081 1.00 38.64 ATOM 3613 C ARG B 74 22.142 10.351 83.297 1.00 55.84 ATOM 3613 C ARG B 74 22.142 10.351 83.297 1.00 55.84 ATOM 3613 C ARG B 74 22.142 10.351 83.297 1.00 55.84 ATOM 3615 C ARG B 74 22.142 10.351 83.297 1.00 55.84 ATOM 3612 C ARG B 74 22.142 10.351 83.297 1.00 34.37 ATOM 3613 C ARG B 74 22.142 10.351 83.297 1.00 35.69 ATOM 3622 C ARG B 74 22.142 10.351 83.297 1.00 55.84 ATOM 3622 C ARG B 74 22.142 10.351 83.297 1.00 35.92 ATOM 3622 C ARG B 74 22.142 10.351 83.297 1.00 35.92 ATOM 3622 C ARG B 74 22.142 10.351 83.297 1.00 35.69 ATOM 3622 C ARG B 74 22.142 10.351 83.297 1.00 35.69 ATOM 3622 C ARG B 74 22.142 10.351 83.297 1.00 35.84 ATOM 3622 C ARG B 74 22.142 10.351 83.297 1.00		3592	OE1 G	LU E	3 71					
ATOM 3594 C GLU B 71 26.559 6.209 84.887 1.00 29.15 ATOM 3595 O GLU B 71 26.115 7.341 85.481 1.00 27.76 ATOM 3596 N ALA B .72 27.671 5.781 85.481 1.00 27.76 ATOM 3597 CA ALA B .72 28.454 6.662 86.340 1.00 27.58 ATOM 3599 C ALA B .72 28.924 7.886 85.563 1.00 28.07 ATOM 3500 O ALA B .72 28.895 8.999 86.909 1.00 23.24 ATOM 3601 N GLU B .73 29.356 7.684 84.322 1.00 26.40 ATOM 3601 N GLU B .73 29.356 7.684 84.322 1.00 26.40 ATOM 3603 CB GLU B .73 30.658 8.314 82.325 1.00 29.06 ATOM 3604 CG GLU B .73 31.162 9.466 81.443 1.00 31.00 ATOM 3605 CD GLU B .73 31.938 9.009 80.216 1.00 34.37 ATOM 3606 OEI GLU B .73 31.938 9.009 80.216 1.00 34.37 ATOM 3608 C GLU B .73 33.059 8.461 80.356 1.00 28.41 ATOM 3609 O GLU B .73 31.419 9.203 79.100 1.00 30.59 ATOM 3608 C GLU B .73 28.8744 9.734 83.045 1.00 31.92 ATOM 3601 N ARG B .74 27.633 9.186 82.570 1.00 33.53 ATOM 3601 N ARG B .74 26.583 10.067 82.081 1.00 35.69 ATOM 3610 N ARG B .74 26.583 10.067 82.081 1.00 35.69 ATOM 3611 CA ARG B .74 26.583 10.067 82.081 1.00 33.53 ATOM 3612 CB ARG B .74 26.583 10.067 82.081 1.00 38.64 ATOM 3613 CG ARG B .74 24.448 8.706 82.365 1.00 55.58 ATOM 3616 CZ ARG B .74 24.448 8.706 82.365 1.00 58.93 ATOM 3617 NHI ARG B .74 22.142 10.051 83.297 1.00 55.58 ATOM 3618 NE ARG B .74 22.142 10.351 83.297 1.00 58.93 ATOM 3618 NE ARG B .74 22.142 10.351 83.297 1.00 58.93 ATOM 3618 NE ARG B .74 22.142 10.351 83.297 1.00 58.93 ATOM 3618 NE ARG B .74 22.142 10.351 83.297 1.00 58.93 ATOM 3618 NE ARG B .74 22.142 10.051 89.10 84.192 1.00 35.69 ATOM 3618 NE ARG B .74 22.142 10.05 84.386 1.00 35.45 ATOM 3618 NE ARG B .74 22.142 10.351 83.297 1.00 33.62 ATOM 3620 C ARG B .74 22.142 10.351 83.297 1.00 33.62 ATOM 3621 C BR B .75 25.581 9.228 87.123 1.00 34.47 ATOM 3622 CA SER B .75 25.581 9.228 87.123 1.00 34.33 ATOM 3623 CB GLN B .76 28.818 13.572 86.241 1.00 33.45 ATOM 3624 CG SER B .75 25.581 9.288 87.123 1.00 33.45 ATOM 3625 C GLN B .76 31.124 13.681 85.248 1.00 35.43 ATOM 3623 CG GLN B .76 31.124 13.681 85.248 1.00 35.43 ATOM 3634 C GLN B .76 31.124 13.681 85.248	,	3593	OE2 G	LU E	3 71					
ATOM 3595 O GLU B 71										
ATOM 3596 N ALA B 72										
ATOM 3597 CA ALA B 72 28.454 6.662 86.340 1.00 27.58 ATOM 3598 CB ALA B 72 28.924 7.886 85.563 1.00 28.07 ATOM 3600 O ALA B 72 28.8924 7.886 85.563 1.00 28.07 ATOM 3601 N GLU B 73 29.356 7.684 84.322 1.00 26.40 ATOM 3602 CA GLU B 73 29.356 7.684 84.322 1.00 26.40 ATOM 3602 CA GLU B 73 30.658 8.314 82.325 1.00 29.06 ATOM 3603 CB GLU B 73 30.658 8.314 82.325 1.00 29.48 ATOM 3605 CD GLU B 73 31.162 9.466 81.443 1.00 31.00 31.07 ATOM 3605 CD GLU B 73 31.938 9.009 80.216 1.00 34.37 ATOM 3606 OEI GLU B 73 31.938 9.009 80.216 1.00 34.37 ATOM 3606 OEI GLU B 73 31.419 9.203 79.100 1.00 30.59 ATOM 3608 C GLU B 73 31.419 9.203 79.100 1.00 30.59 ATOM 3608 C GLU B 73 28.894 10.951 83.104 1.00 31.02 ATOM 3608 C GLU B 73 28.894 10.951 83.104 1.00 35.69 ATOM 3611 CA ARG B 74 26.583 10.067 82.081 1.00 38.64 ATOM 3612 CB ARG B 74 26.583 10.067 82.081 1.00 38.64 ATOM 3613 CG ARG B 74 26.583 10.067 82.081 1.00 38.64 ATOM 3614 CD ARG B 74 22.4448 8.706 82.363 1.00 46.67 ATOM 3615 NE ARG B 74 22.142 10.351 83.297 1.00 58.93 ATOM 3616 CC ARG B 74 22.076 8.153 82.594 1.00 55.58 ATOM 3617 NH ARG B 74 22.142 10.351 83.297 1.00 58.93 ATOM 3620 C ARG B 74 22.076 8.153 82.594 1.00 55.58 ATOM 3617 NH ARG B 74 22.142 10.351 83.297 1.00 58.93 ATOM 3620 C ARG B 74 22.076 8.153 82.594 1.00 55.58 ATOM 3620 C ARG B 74 22.076 8.153 82.597 1.00 58.93 ATOM 3620 C ARG B 74 22.076 8.153 82.597 1.00 58.93 ATOM 3620 C ARG B 74 22.142 10.351 83.297 1.00 58.93 ATOM 3620 C ARG B 74 22.142 10.351 83.297 1.00 58.93 ATOM 3620 C ARG B 74 22.142 10.351 83.297 1.00 33.45 ATOM 3620 C ARG B 74 22.142 10.351 83.297 1.00 33.45 ATOM 3620 C ARG B 75 25.588 75 25.581 9.228 87.123 1.00 34.37 ATOM 3620 C ARG B 74 22.142 10.351 83.297 1.00 33.45 ATOM 3620 C ARG B 75 24.599 10.038 86.510 1.00 34.37 ATOM 3620 C ARG B 75 24.599 10.308 86.510 1.00 33.45 ATOM 3623 CG GLN B 76 28.818 13.052 28.66.241 1.00 30.11 ATOM 3633 CG GLN B 76 31.052 12.546 84.761 1.00 39.13 ATOM 3631 C G GLN B 76 31.052 12.546 84.761 1.00 39.31 ATOM 3633 CG GLN B 76 31.052 12.546 84.761 1.00 39.31 ATOM 3										
ATOM 3598 CB ALA B 72										
ATOM 3600 O ALA B 72 28.924 7.886 85.563 1.00 28.07 ATOM 3601 N GLU B 72 28.895 8.999 86.079 1.00 23.22 ATOM 3601 N GLU B 73 29.356 7.684 84.322 1.00 26.40 ATOM 3602 CA GLU B 73 30.658 8.314 82.325 1.00 29.06 ATOM 3603 CB GLU B 73 30.658 8.314 82.325 1.00 29.08 ATOM 3605 CD GLU B 73 31.938 9.009 80.216 1.00 34.37 ATOM 3606 OEI GLU B 73 33.059 8.461 80.356 1.00 28.41 ATOM 3606 OEI GLU B 73 33.059 8.461 80.356 1.00 28.41 ATOM 3607 OE2 GLU B 73 33.059 8.461 80.356 1.00 34.37 ATOM 3608 C GLU B 73 28.894 10.951 83.104 1.00 35.69 ATOM 3600 N ARG B 74 26.583 10.067 82.081 1.00 38.64 ATOM 3611 CA ARG B 74 26.583 10.067 82.081 1.00 38.64 ATOM 3612 CB ARG B 74 25.456 9.280 81.403 1.00 39.90 ATOM 3613 CG ARG B 74 22.076 8.153 82.594 1.00 55.58 ATOM 3616 CZ ARG B 74 22.076 8.153 82.297 1.00 58.93 ATOM 3617 NH1 ARG B 74 22.076 8.153 82.297 1.00 58.93 ATOM 3620 C ARG B 74 22.006 8.153 82.297 1.00 58.93 ATOM 3621 N SER B 75 25.798 12.107 83.048 1.00 33.45 ATOM 3622 CA SER B 75 25.581 9.228 87.120 1.00 33.45 ATOM 3623 CB SER B 75 25.581 9.228 87.120 1.00 33.45 ATOM 3624 OG SER B 75 25.581 9.228 87.120 1.00 33.45 ATOM 3625 C SER B 75 25.581 9.228 87.120 1.00 33.45 ATOM 3626 C SER B 75 25.581 9.228 87.120 1.00 33.45 ATOM 3627 N GLN B 76 28.818 13.572 86.241 1.00 30.11 ATOM 3631 CD GLN B 76 31.052 12.546 84.761 1.00 39.13 ATOM 3632 OEI GLN B 76 31.052 12.546 84.761 1.00 39.13 ATOM 3633 NEZ GLN B 76 31.052 12.546 84.761 1.00 39.13 ATOM 3635 O GLN B 76 31.052 12.546 84.761 1.00 39.13 ATOM 3631 CD GLN B 76 31.052 12.546 84.761 1.00 39.13 ATOM 3633 OCG GLN B 76 31.1995 14 583 84.827 1.00 47.88 ATOM 3635 O GLN B 76 31.1995 14 583 84.827 1.00 47.88 ATOM 3635 O GLN B 76 31.1995 14 583 84.827 1.00 37.88 ATOM 3635 O GLN B 76 31.1995 14 583 84.827 1.00 47.88 ATOM 3635 O GLN B 76 31.1995 14 583 84.827 1.00 47.88 ATOM 3635 O GLN B 76 31.1995 14 583 84.827 1.00 47.88 ATOM 3635 O GLN B 76 31.1995 14 583 84.827 1.00 47.93										
ATOM 3600 O ALA B 72										
ATOM 3601 N GLU B 73										
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ATOM 3629 CB GLN B 76 28.818 13.572 86.241 1.00 30.11 ATOM 3630 CG GLN B 76 30.216 14.112 86.390 1.00 39.13 ATOM 3631 CD GLN B 76 31.124 13.681 85.248 1.00 33.54 ATOM 3632 OE1 GLN B 76 31.052 12.546 84.761 1.00 29.21 ATOM 3633 NE2 GLN B 76 31.995 14 583 84.827 1.00 40.93 ATOM 3634 C GLN B 76 28.624 11 892 88.079 1.00 37.88 ATOM 3635 O GLN B 76 28.901 12 808 88.858 1.00 32.74	ATOM	3627	N GL	N B	76		27.584	11.473	85.922	1.00 33.25
ATOM 3629 CB GLN B 76 28.818 13.572 86.241 1.00 30.11 ATOM 3630 CG GLN B 76 30.216 14.112 86.390 1.00 39.13 ATOM 3631 CD GLN B 76 31.124 13.681 85.248 1.00 33.54 ATOM 3632 OE1 GLN B 76 31.052 12.546 84.761 1.00 29.21 ATOM 3633 NE2 GLN B 76 31.995 14 583 84.827 1.00 40.93 ATOM 3634 C GLN B 76 28.624 11 892 88.079 1.00 37.88 ATOM 3635 O GLN B 76 28.901 12 808 88.858 1.00 32.74	ATOM	3628	CA GL	N B	76		28.739	12.082	86.565	1.00 35.61
ATOM 3630 CG GLN B 76 30.216 14.112 86.390 1.00 39.13 ATOM 3631 CD GLN B 76 31.124 13.681 85.248 1.00 33.54 ATOM 3632 OE1 GLN B 76 31.052 12.546 84.761 1.00 29.21 ATOM 3633 NE2 GLN B 76 31.995 14.583 84.827 1.00 40.93 ATOM 3634 C GLN B 76 28.624 11.892 88.079 1.00 37.88 ATOM 3635 O GLN B 76 28.901 12.808 88.858 1.00 32.74		3629	CB GL	N B	76		28.818	13.572	86.241	
ATOM 3631 CD GLN B 76 31.124 13.681 85.248 1.00 33.54 ATOM 3632 OE1 GLN B 76 31.052 12.546 84.761 1.00 29.21 ATOM 3633 NE2 GLN B 76 31.995 14 583 84.827 1.00 40.93 ATOM 3634 C GLN B 76 28.624 11 892 88.079 1.00 37.88 ATOM 3635 O GLN B 76 28.901 12 808 88.858 1.00 32.74	ATOM	3630	CG GL	N B	76		30.216	14.112	86.390	1.00 39.13
ATOM 3633 NE2 GLN B 76 31.995 14 583 84.827 1.00 40.93 ATOM 3634 C GLN B 76 28.624 11 892 88.079 1.00 37.88 ATOM 3635 O GLN B 76 28.901 12 908 88.858 1.00 32.74	ATOM	3631	CD GL	NB	76		31.124	13.681	85.248	1.00 33.54
ATOM 3634 C GLN B 76 28.624 11 892 88.079 1.00 37.88 ATOM 3635 O GLN B 76 28.901 12 908 88.858 1.00 32.74	ATOM	3632	OE1 GL	N B	76		31.052	12.546	84.761	1.00 29.21
ATOM 3635 O GLN B 76 28.901 12 808 88.858 1.00 32.74	MOTA	3633	NE2 GL	N B	76		31.995	14 583	84.827	1.00 40.93
	ATOM	3634	C GL	B	76		28.624	11 892	88.079	
ATOM 3636 N SER B 77 28.209 10.697 88.488 1.00 34.72	MOTA	3635						12 908	88.858	
	MOTA	3636	N SE	R B	77		28.209	10.697	88.488	1.00 34.72
ATOM 3637 CA SER B 77 28.047 10.382 89.901 1.00 37.07										1.00 37.07
ATOM 3638 CB SER B 77 26.635 10.738 90.371 1.00 39.61		3638	CB SEI	RB	77		26.635	10.738	90.371	1.00 39.61
ATOM 3639 CG SER B 77 25.678 9.941 89.688 1.00 39.03		3639	OG SEI	RB	7 7		25.678	9.941	89.688	1.00 39.03
ATOM 3640 C SER B 77 28.265 8.897 90.112 1.00 35.95		3640	C SEI	RB	77		28.265		90.112	1.00 35.95
ATOM 3641 O SER B 77 28.177 8.108 89.173 1.00 36.60		3641	O SER	B	77		28.177	8.108	89.173	1.00 36.60
ATOM 3642 N VAL B 78 28.528 8.518 91.355 1.00 33.03		3642	N VAI	, B	78		28.528		91.355	1.00 33.03
ATOM 3643 CA VAL B 78 28.753 7.124 91.685 1.00 33.41		3643			78				91.685	1.00 33.41
ATOM 3644 CB VAL B 78 29.742 6.979 92.848 1.00 36.91		3644	CB VAI	, В	78				92.848	1.00 36.91
ATOM 3645 CG1 VAL B 78 29.955 5.499 93.163 1.00 34.37			CG1 VAI	, B	78				93.163	
ATOM 3646 CG2 VAL B 78 31.055 7.658 92.496 1.00 34.19		3646	CG2 VAI	, в	78		31.055			
ATOM 3647 C VAL B 78 27.461 6.431 92.082 1.00 34.93		3647			78		27.461			1.00 34.93
ATOM 3648 0 VAL B 78 26.897 6.703 93.143 1.00 28.25		3648	IAV C	, В			26.897	6.703	93.143	
ATOM 3649 N PRO B 79 26.971 5.521 91.228 1.00 36.73		3649	N PRO	В			26.971			1.00 36.73
ATOM 3650 CD PRO B 79 27.532 5.114 89.930 1.00 37.44		3650	CD PRO	В			27.532	5.114		
ATOM 3651 CA PRO B 79 25.738 4.779 91.493 1.00 38.33			CA PRO	В			25.738			1.00 38.33
ATOM 3652 CB PRO B 79 25.668 3.826 90.301 1.00 38.68			CB PRO	В		•	25.668			
ATOM 3653 CG PRO B 79 26.293 4.664 89.201 1.00 37.41		3653	CG PRO	В	79		26.293	4.664	89.201	1.00 37.41

			220						
ATOM			PRO		25.788		92.834	1.00	36.92
MOTA					26.85	3.648	93.298	1.00	33.03
ATOM	1 365	6 N	LYS	B 80	24.623	3.881			38.43
ATOM	365	7 C	A LYS		24.482				
ATOM								1.00	39.73
				-	23.003				43.33
ATOM					22.679	2:129	96.262	1.00	44.60
MOTA	366	a ci	LYS	B 80	21.198	1.742			48.09
ATOM	366	1 CE	LYS	B 80	20.805				
ATOM		_							50.12
					20.932	1.890		1.00	53.16
ATOM	366	3 C	LYS		25.315	1.928	94.854	1.00	40.35
ATOM	3664	4 0	LYS	B 80	- 25.181				36.67
MOTA	3665	5 N	GLY		26.173				
									38.26
ATOM					26.996				34.69
ATOM	366	7 C	GLY :	B 81	28.066	0.407	95.071	1.00	34.63
MOTA	3668	3 0	GLY :	B 81	28.861				33.92
ATOM		N	ALA :		28.100				
									31.26
ATOM					. 29.082			1.00	34.88
MOTA	3671	CB	ALA I	B 82.	28.755	1.848	91.751	1.00	23.13
MOTA	3672	: C	ALA 1	82	30.517	1.223	93.405		36.85
ATOM	3673		ALA I		31.461	0.580			
							92.945		32.17
ATOM	3674		ARG I		30.677	2.168	94.323	1.00	36.52
ATOM	3675	CA	ARG I	83	31.994	2.522	94.830	1.00	38.75
MOTA	3676	CB	ARG E	83	31.865	3.616			40.24
ATOM	3677		ARG E		33.187				
						4.180	96.330		49.12
MOTA	3678		ARG E		33.015	5.239	97.404	1.00	53.26
ATOM	3679	NE	ARG E	83	34.240	6.010	97.624	1.00	59.30
ATOM	3680	CZ	ARG E	83	35.437	5.486	97.883		61.56
ATOM	3681	NH:	L ARG E		35.598	4.170	97.958		
ATOM	3682								63.53
					36.479	6.285	98.073	1.00	62.02
ATOM -		С	ARG B		32.719	1.326	95.426	1.00	37.75
MOTA	3684	0	ARG B	83	33.893	1.094	95.146	1.00	37.18
MOTA	3685	N	GLU B	84	32.011	0.564	96.249		35.29
ATOM	3686	CA	GLU B		32.581		,		
						-0.609	96.898		35.29
ATOM	3687	CB	GLU B		31.876	-0.855	98.236	1.00	40.14
MOTA	3688	CG	GLU B	84	30.443	-0.383	98.240		46.30
ATOM	3689	CD	GLU B	84	30.356	1.132	98.293		48.30
ATOM	3690	OE1	GLU B	84	29.339	1.690			
		0E2					97.834		43.07
MOTA	3691			84	31.306	1.762	98.814	1.00	50.07
MOTA	3692	С	GLU B	84	32.527	-1.880	96.055	1.00	32.90
MOTA	3693	0	GLU B	84	33.371	-2.765	96.193	1.00	28.68
ATOM	3694	Ν.	LYS B	85	31.533	-1.984	95.187		27.12
ATOM	3695	CA	LYS B	85					
					31.412	-3.177	94.361		30.46
ATOM	3696	CB	LYS B	85	29.950	-3.401	93.967	1.00	30.01
ATOM	3697	CG	LYS B	85	29.717	-4.643	93.117	1.00	28.40
ATOM	3698	CD	LYS B	85	28.234	-4.807	92.775		32.87
A. OM	3699	CE	LYS'B	85	28.000	-6.048	91.928		34.15
A.OM	3700	NZ	LYS B	85					
					26.582	-6:186	91.507		35.34
MOLA	3701	С	LYS B	85	32.267	-3.096	93.101	1.00	28.98
ATOM	3702	0	LYS B	85	32.817	-4.098	92.652	1.00	24.69
ATOM	3703	N	TYR B	86	32.391	-1.896	92.550		27.81
ATOM	3704	CA	TYR B	86	33.141				
						-1.692	91.319		27.56
ATOM	3705	CB	TYR B	86	32.206	-1.050	90.288	1.00	28.88
ATOM ·	3706	CG	TYR B	86	31.008	-1.927	89.951	1.00	31.29
ATOM	3707	CD1	TYR B	86	31.178	-3.137	89.276	1.00	
ATOM	3708	CE1	TYR B	86	30.095		88.965		
						-3.955	88.963	1.00	
MOTA	3709	CD2	TYR B	86	29.713	-1.553	90.315	1.00	28.38
ATOM	3710	CE2	TYR B	86	28.611	-2.370	90.008	1.00	24.19
MOTA	3711	CZ	TYR B	86	28.815	-3.569	89.331	1.00	
ATOM	3712	OH	TYR B	86	27.747	-4.379	89.008		
								1.00	
MOTA	3713	C	TYR B	86	34.422	-0.870	91.489	1.00	24.64
MOTA	3714	0	TYR B	86	35.160	-0.645	90.530	1.00	27.19
MOTA	3715	N	ASN B	87	34.674	-0.418	92.711	1.00	
ATOM	3716		ASN B	87	35.881	0.341	93.032	1.00	
ATOM	3717		ASN B	87	37.105	-0.561	92.866	1.00	
ATOM	3718		asn B	87	38.343	0.019	93.506	1.00	34.72
MOTA	3719	OD1	asn B	87	38.309	0.452	94.659	1.00	

ATOM	3720	ND2	ASN	R	87	39.449	0.012	92.775	1.00 35.86
								92.223	1.00 29.72
ATOM	3721	С	ASN	B	87	36.070	1.622		
MOTA	3722	0	ASN	В	87	37.194	1.998	91.876	1.00 24.01
						34.956	2.282	91.932	1.00 29.43
MOTA	3723	N	ILE	В	88				•
ATOM	3724	CA	ILE	В	88	34.945	3.536	91.196	1.00 30.64
	3725	CB	ILE		88	33.959	3.464	90.027	1.00 37.12
ATOM-									
ATOM	3726	CG2	ILE	В	88	33.821	4.829	89.379	1.00 40.62
	3727	CG1	ILE	R	88	34.421	2.433	89.008	1.00 35.43
MOTA									
MOTA	3728	CD1	ILE	В	88	35.684	2.821	88.324	1.00 41.80
ATOM	3729	С	-ILE	В	88	34.483	4.669	92.118	1.00 31.90
								93.024	1.00 28.86
ATOM	3730	0	ILE	В	88	33.681	4.445		
MOTA	3731	N	GLY	B.	89	34.977	5.881	91.875	1.00 30.36
•					89 .	34.574	7.022	92.686	1.00 29.54
ATOM	3732	CA	GLY						
ATOM	3733	С	GLY	В	89	35.601	7.524	93.685	1.00 31.49
	3734	0	GLY	B	89	35.497	8.652	94.177	1.00 37.26
ATOM								94.005	1.00 30.97
ATOM	3735	N	GLY	В	90	36.583	6.687		
ATOM	3736	CA	GLY	В	90	37.612	7.086	94.949	1.00 31.03
					90	38.655	7.936	94.247	1.00 34.78
ATOM	3737	С	GLY						
ATOM	3738	0	GLY	В	90	38.455	8.344	93.103	1.00 32.73
	3739	N	TYR	B	91	39.772	8.201	94.915	1.00 29.39
MOTA.									1.00 28.15
ATOM	3740	CA	TYR	В	91	40.820	9.023	94.322	
ATOM	3741	CB	TYR	В	91	41.810	9.463	95.405	1.00 27.29
					91	42.609	8.330	96.007	1.00 26.60
MOTA	3742	CG	TYR						
ATOM	3743	CD1	TYR	В	91	43.738	7.823	95.359	1.00 28.55
	3744	CE1	TYR	R	91	44.456	6.762	95.896	1.00 28.75
MOTA							7.741	97.208	1.00 28.35
ATOM	3745	CD2	TYR	В	91	42.219			
MOTA	3746	CE2	TYR	В	91	42.927	6.680	97.751	1.00 27.58
					91	44.043	6.196	97.094	1.00 30.12
ATOM	3747	CZ	TYR						
ATOM	3748	OH	TYR	В	91	44.753	5.154	97.637	1.00 36.59
ATOM	3749	С	TYR	В	91	41.563	8.271	93.226	1.00 29.27
			TYR		91	42.109	8.874	92.308	1.00 25.22
ATOM	3750	0							1.00 28.32
ATOM	3751	N	GLU	В	92	41.568	6.948	93.318	
ATOM	3752	CA	GLU	В	92	42.286	6.124	92.350	1.00 27.06
	3753	CB	GLU		92	42.474	4.726	92.924	1.00 23.35
MOTA								92.221	1.00 29.80
ATOM	3754	CG	GLU	В	92	43.502	3.884		
ATOM	3755	CD	GLU	В	92	43.585	2.500	92.826	1.00 35.34
	3756	OFI	GLU	B	92	42.742	1.645	92.477	1.00 32.15
ATOM							2.278	93.678	1.00 31.61
ATOM	3757	OE2	GLU		92	44.475			
ATOM	3758	С	GLU	В	92	41.594	6.024	90.997	1.00 23.42
	3759	0	GLU		92	42.204	6.260	89.962	1.00 20.47
MOTA							5.677	91.017	1.00 18.85
ATOM	3760	N	ASN		93	40.314			
MOTA	3761	CA	ASN	В	93	39.534	5.509	89.795	1.00 21.96
	3762	CB	ASN	R	93	39.165	4.033	89.664	1.00 23.90
ATOM							3.120	89.943	1.00 24.78
ATOM	3763	CG	ASN		93	40.351			
ATOM	3764	OD1	ASN	В	93	41.362	3.160	89.239	1.00 22.35
	_	ND2	ASN	R	93	40.240	2.311	90.987	1.00 13.35
ATOM	3765							89.944	1.00 25.16
MOTA	3766	C	asn	B	93	38.285	6.362		
ATOM	3767	0	ASN	В	93	37.183	5.843	90.121	1.00 20.91
			PRO		94	38.449	7.693	89.887	1.00 26.37
ATCM	3768	N							1.00 19.35
MOTA	3769	CD	PRO	В	94	39.738	8.389	89.716	
	3770	CA	PRO	R	94	37.373	8.676	90.024	1.00 24.59
MOTA								90.200	1.00 25.95
MOTA	3771	CB	PRO		94	38.147	9.972		1.00 25.55
MOTA	3772	CG	PRO	В	94	39.297	9.740	89.223	1.00 22.60
					94	36.384	8.777	88.873	1.00 28.74
MOTA	3773	С	PRO						
MOTA	3774	0	PRO	В	94	36.562	8.176	87.808	1.00 25.77
	3775	N	VAL	В	95	35.332	9.553	89.112	1.00 27.14
Mota								88.103	1.00 25.94
MOTA	3776	CA	VAL		95	34.317	9.812		1.00 23.75
ATOM	3777	CB	VAL	В	95	33.035	10.393	88.742	
	3778	CG1	VAL	В	95	32.067	10.855	87.662	1.00 26.34
ATOM							9.346	89.622	1.00 27.59
ATOM	3779	CG2	VAL		95	32.376			
ATOM	3780	С	VAL		95	34.912	10.861	87.175	1.00 25.69
	3781	0	VAL	В	95	35.564	11.793	87.641	1.00 25.25
ATOM	-		SER		96	34.708	10.699	85.871	1.00 28.02
MOTA	3782	N						84.868	1.00 24.82
ATOM	3783	CA	SER		96	35.199	11.647		
	3784	CB	SER	В	96	36.729	11.705	.84.850	1.00 25.90
ATOM	3,05	OG	SER		96	37.274	10.548	84.229	1.00 23.99
MOTA	3785	0.3	٠٠٠٠	_	- 0	5	10.555	_	_

ATOM	3786	C	SER B	96		34.726	11.127	83.519	1.00	26.22
ATOM	3787	0	SER B	96		33.943	10.174	83.462	1.00	23.57
ATOM	3788	N	TYR B	97		35.195	11.744	82.438	1.00	22.83
ATOM	3789		TYR B	97		34.818	11.279	81.110		28.59
ATOM	3790		TYR B	97		34.536	12.452			
								80.165		31.45
ATOM	3791	CG	TYR B	97		33.279	13.203	80.548		35.09
ATOM	3792	CD		97		33.316	14.239	81.480		32.87
ATOM	3793	CE	1 TYR B	97		32.148	14.863	81.911	1.00	37.73
ATOM	3794	CD	2 TYR B	97		32.036	12.812	80.049	1.00	34.85
ATOM	3795	CE:	2 TYR B	97		30.858	13.430	80.475		38.61
ATOM	3796	CZ	TYR B	97		30.924	14.453	81.408		39.45
ATOM	3797	ОН	TYR B	97		29.768	15.047	81.852		35.36
ATOM	3798	C	TYR B	97		35.883	10.354	80.534		
	3799									28.93
ATOM		0	TYR B	97		35.859	9.992	79.358		28.26
ATOM	3800	N	ALA B	98		36.822	9.968	81.385		29.09
ATOM	3801	CA	ALA B	98		37.866	9.044	80.980	1.00	26.88
ATOM	3802	CB	ALA B	98.		39.167	9.369	81.692		27.99
MOTA	3803	С	ALA B	98		37.395	7.657	81.382	1.00	22.53
ATOM	3804	0	ALA B	98		37.721	6.675	80.722		21.98
ATOM	3805	N	MET B	99		36.603	7.595	82.453		23.51
ATOM	3806	CA	MET B	99		36.106	6.326	82.986		26.36
ATOM	3807	СВ	MET B	99		35.179	6.568	84.185		24.05
	3808	CG	MET B	99		33.822	7.188	83.875	1.00	
ATOM										28.37
ATOM	3809	SD	MET B	99		32.966	7.704	85.406		27.91
ATOM	3810	CE	MET B	99		33.106	6.227	86.409		22.12
ATOM	3811	C	MET B	99		35.430	5.435	81.953		25.76
ATOM	3812	0	MET B	9 9		35.544	4.212	82.031	1.00	26.11
ATOM	3813	N	PHE B 1	.00		34.724	6.027	80.992	1.00	22.17
MOTA	3814	CA	PHE B 1	.00		34.107	5.222	79.940	1.00	22.35
ATOM	3815	CB	PHE B 1	.00		32,582	5.133	80.088	1.00	22.01
ATOM	3816	CG	PHE B 1	.00		31.947	4.254	79.038	1.00	24.22
ATOM	3817	CD1	PHE B 1	.00		32.143	2.872	79.061	1.00	26.61
ATOM	3818	CD2		00		31.280	4.813	77.953		21.22
ATOM	3819			00		31.691	2.059	78.012		26.91
ATOM	3820	CE2		00		30.825	4.010	76.894		24.80
ATOM	3821	CZ		00		31.033	2.632	76.924		24.85
ATOM	3822	c		00		34.425	5.695	78.514		24.86
	3823	ŏ		00		34.922	4.920	77.694		21.40
MOTA										
MOTA	3824	N		01		34.131	6.957	78.204	1.00	
ATOM	3825	CA	THR B 1			34.390	7.469	76.854		24.54
ATOM	3826	CB		01		33.914	8.926	76.708		24.46
MOTA	3827	OG1				32.504	8.985	76.953	1.00	27.64
ATOM	3828	CG2	THR B 1	01	3	34.191	9.445	75.297	1.00	22.19
ATOM	3829	С	THR B 1	01	3	35.872	7.387	76.483	1.00	25.26
MOTA	3830	0	THR B 1	01	3	36.231	6.856	75.430	1.00	25.47
ATOM	3831	· N	GLY B 1	02	3	6.725	7.916	77.350	1.00	23.74
ATOM	3832	CA	GLY B 1	02	3	8.153	7.867	77.096	1.00	24.53
ATOM	3833	C	GLY B 1	02		8.657	6.434	77.046	1.00	24.06
ATOM	3834	O	GLY B 1			9.346	6.045	76.100		22.53
ATOM	3835	N	SER B 1			8.316	5.651	78.067		22.02
	3836	CA	SER B 1			8.730	4.253	78.146	1 00	20.45
ATOM					2	0.750				
ATOM	3837	CB	SER B 10			8.193	3.613	79.427		25.21
MOTA	3838	OG	SER B 10			8.820	4.166	80.567		26.48
MOTA	3839	С	SER B 10			8.268	3.446	76.938		20.53
MOTA	3840	0	SER B 10)3	3	9.034	2.669	76.372		16.82
MOTA	3841	N	SER B 10) 4	3	7.014	3.642	76.542	1.00	17.11
MOTA	3842	CA	SER B 10)4	3	6.462	2.937	75.395	1.00	23.32
ATOM	3843	CB	SER B 10	14	3	4.980	3.289	75.228	1.00	22.93 .
MOTA	3844	OG	SER B 10			4.424	2.557	74.161		24.75
ATOM	3845	C	SER B 10			7.221	3.288	74.116		21.97
	3846		SER B 10			7.451	2.434	73.256		22.83
ATOM			LEU B 10			7.619	4.549	73.230		23.00
MOTA	3847		LEU B 10			8.354	5.007	72.825		25.12
ATCM	3848									29.25
ATOM	3849		LEU B 10			8.443	6.536	72.859	_	
MOTA	3850	CG	LEU B 10	13		8.702	7.289	71.553		34.27
ATCM	3851	CDI	LEU B 10	, 5	3	7.662	6.888	70.512	1.00	31.51

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			•			
MOTA	3852	CD2 LEU B 105	5 38.62	9 8.802	71.819	1.00 34.30
ATOM	3853				72.813	1.00 27.39
MOTA	3854				71.765	1.00 27.39
	3855					
MOTA					73.988	1.00 23.62
ATOM	3856				74.115	1.00 22.73
MOTA	3857				75.529	1.00 17.46
ATOM	3858			9 2.189	73.846	1.00 22.77
MOTA	3859	O ALA B 106	42.58	3 1.597	73.316	1.00 24.98
ATOM '	3860	N THR B 107	7 40.52	3 1.567	74.224	1.00 23.54
ATOM	3861	CA THR B 107	40.35	5 0.132	74.033	1.00 19.76
MOTA	3862				74.947	1.00 22.85
ATOM	3863				76.306	1.00 16.29
ATOM	3864	CG2 THR B 107			74.787	1.00 17.21
ATOM	3865				72.571	1.00 23.47
	3866	O THR B 107			72.001	
ATOM						1.00 19.25
MOTA	3867				71.959	1.00 24.87
ATOM	3868	CA GLY B 108			70.560	1.00 21.32
MOTA	3869	C GLY B 108			69.757	1.00 22.01
ATOM	3870	O GLY B 108			68.761	1.00 20.46
MOTA	3871	N SER B 109			70.197	1.00 19.89
MOTA	3872	CA SER B 109			69.499	1.00 19.37
MOTA	3873	CB SER B 109	42.993	3 2.956	70.107	1.00 18.28
ATOM	3874	OG SER B 109	42.250	4.145	69.882	1.00 21.29
MOTA	3875	C SER B 109	43.168	0.513	69.542	1.00 20.70
ATOM	3876	O SER B 109	43.940	0.261	68.617	1.00 20.69
ATOM	3877	N THR B 110			70.616	1.00 20.54
ATOM	3878	CA THR B 110			70.729	1.00 19.98
ATOM	3879	CB THR B 110			72.158	1.00 20.63
ATOM	3880	OG1 THR B 110			73.007	1.00 20.72
ATOM	3881	CG2 THR B 110			72.188	1.00 20.05
ATOM	3882	C THR B 110	43.333		69.738	1.00 21.61
ATOM	3883	O THR B 110	44.115		69.127	1.00 18.11
	3884	N VAL B 111	42.012		69.567	1.00 18.11
ATOM	3885	CA VAL B 111	41.432			
ATOM		•			68.608	1.00 20.36
ATOM	3886		39.886		68.677	1.00 23.94
ATOM	3887	CG1 VAL B 111	39.324		67.619	1.00 24.37
MOTA	3888	CG2 VAL B 111	39.426		70.063	1.00 21.60
MOTA	3889	C VAL B 111	41.872		67.197	1.00 20.35
MOTA	3890	O VAL B 111	42.146		66.362	1.00 23.29
ATOM .	3891	N GLN B 112	41.953		66.937	1.00 22.07
ATOM	3892	CA GLN B 112	42.367		65.617	1.00 22.34
ATOM	3893	CB GLN B 112	42.199		65.513	1.00 24.54
MOTA	3894	CG GLN B 112	40.810		65.843	1.00 20.63
ATOM	3895	CD GLN B 112	40.700		65.742	1.00 21.19
MOTA	3896	OE1 GLN B 112	40.664	2.794	64.645	1.00 26.73
ATOM	389 7	NE2 GLN B 112	40.667	2.905	66.886	1.00 18.73
ATOM	3898	C GLN B 112	43.826	-1.635	65.363	1.00 23. 1
ATOM	389 9	O GLN B 112	44.195	-2.020	64.257	1.00 15.79
MOTA	3900	N ALA B 113	44.660	-1.476	66.389	1.00 20.60
ATOM	3901	CA ALA B 113	46.070		66.249	1.00 18.02
MOTA	3902	CB ALA B 113	46.794			1.00 20.84
ATOM	3903	C ALA B 113	46.170		65.863	1.00 23.78
MOTA	3904	O ALA B 113	46.982		65.023	1.00 19.83
	. 3905	N ILE B 114	45.331		66.477	1.00 21.45
	3906	CA ILE B 114	45.344		66.168	1.00 24.26
ATOM	3907	CB ILE B 114	44.507		67.191	1.00 20.72
ATOM					66.800	
MOTA	3908	CG2 ILE B 114 CG1 ILE B 114	44.476			1.00 21.92
ATOM	3909		45.116	-6.144	68.593	1.00 24.32
ATOM	3910	CD1 ILE B 114	44.364	-6.872	69.694	1.00 19.01
ATOM	3911	C ILE B 114	44.808	-5.765	ύ4.753	1.00 26.75
ATOM	3912	0 ILE B 114	45.305	-6.640	64.032	1.00 20.18
ATOM	3913	N GLU B 115	43.792	-5.009	64.347	1.00 24.69
ATCM	3914	CA GLU B 115	43.243	-5.198	63.005	1.00 29.26
ATOM	3915	CB GLU B 115	42.043	-4.278	62.770	1.00 29.07
ATCM	3916 ·	CG GLU B 115	40.940	-4.421	63.800	1.00 32.31
ATOM	3917	CD GLU B 115	39.757	-3.519	63.516	1.00 38.14

MOTA	3918	OE.	1 GLU I	B 115	39.98	0 -2.374	63.072	1.00 40.63
ATOM	3919				38.60			
			_			-		1.00 39.86
ATOM	3920		GLU I		44.33	4 -4.906	61.974	1.00 31.52
ATOM	3921	0	GLU I	3 115	44.44	4 -5.603	60.964	1.00 26.43
ATOM	3922	N	GLU F	3 116	45.14	1 -3.879		1.00 26.04
ATOM	3923		GLU F		46.22			
								1.00 26.21
ATOM	392,4	CB	_GLU E	3 116	46.90		61.775	1.00 23.21
MOTA	3925	CG	GLU E	3 116 .	46.05	5 -0.983	61.601	1.00 23.82
MOTA	3926	CD	GLU E	3.116	45.57		60.163	1.00 31.43
	3927							
ATOM			1 GLU E		46.42			1.00 26.45
MOTA	3928	OE:	2 GLU E	3 116	44.34	9 -0.771	59.945	1.00 24.59
MOTA	3929	С	GLU E	3 116	47.25	6 -4.644	61.243	1.00 28.60
MOTA	3930	0	GLU E	3 116	47.85		60.189	1.00 25.01
	3931	N						
ATOM			PHE B		47.47		62.363	1.00 26.22
ATOM	3932	CA			48.42	1 -6.425	62.400	1.00 28.05
ATOM	3933	CB	PHE B	3 117	48.51	6 -7.007	63.805	1.00 32.15
MOTA	3934	CĠ	PHE B	117	49.27		63.869	1.00 33.88
ATOM	3935		PHE B					
					50.65		63.713	1.00 33.52
ATOM	3936	CD2			48.60	4 -9.502	64.054	1.00 32.83
ATOM	3937	CE1	PHE B	117	51.35	6 -9.521	63.740	1.00 31.67
MOTA	3938	CE2	PHE B	117		4 -10.710	64.082	1.00 35.69
ATOM	3939	CZ	PHE B			4 -10.717		
							63.926	1.00 36.72
MOTA	3940	C	PHE B		47.92		61.456	1.00 26.43
ATOM	3941	0	PHE B	117	48.68	9 -8.061	60.669	1.00 27.61
ATOM	3942	N	LEU B	118	46.64	2 -7.809	61.551	1.00 23.59
ATOM	3943	CA	LEU B		46.04		60.705	1.00 29.15
ATOM	3944	CB	LEU B		44.58		61.099	1.00 28.78
ATOM	3945	CG	LEU B		44.37	5 -9.478	62.557	1.00 35.24
ATOM	3946	CD1	LEU B	118	42.898	3 -9.763	62.788	1.00 31.92
ATOM	3947		LEU B			5 -10.723	62.856	1.00 33.40
ATOM	3948	C	LEU · B		46.15		59.236	1.00 30.15
MOTA	3949	0	LEU B	118	46.350	9.276	58.379	1.00 27.04
ATOM	3950	N	LYS B	119	46.035	7.128	58.947	1.00 27.96
ATOM	3951	CA	LYS B		46.12		57.569	1.00 26.69
ATOM	3952	СВ	LYS B		45.470			
							57.412	1.00 23.94
MOTA	3953	CG	LYS B		43.998		57.795	1.00 24.41
ATOM	3954	CD	LYS B	119	43.327	7 -3.970	57.350	1.00 27.53
ATOM	3955	CE	LYS B	119	44.024	-2.739	57.886	1.00 33.13
ATOM	3956	NZ	LYS B		43.371		57.428	1.00 27.75
_								
MOTA	3957	C	LYS B		47.577		57.101	1.00 29.12
ATOM	3958	0	LYS B	119	47.864	-6.160	55.984	1.00 35.25
ATOM	3959	N	GLY B	120	48.493	-7.034	57.958	1.00 30.25
MOTA	3960	CA	GLY B	120	49.896		57.585	1.00 28.38
ATOM	3961	C	GLY B		50.642		57.861	
								1.00 27.91
MOTA	3962	0	GLY B		51.775			1.00 22.25
ATOM	3963	N	ASN B	:21	50.024	-4.836	58.600	1.00 25.42
MOTA	3964	CA	ASN B	. 21	50.695	-3.587	58.919	1.00 29.49
MOTA	3965	CB	ASN B		49.758		58.727	1.00 30.07
	3966				49.201			
ATOM		CG	ASN B				57.325	1.00 32.25
ATOM	3967	OD1	ASN B		49.924	-2.491	56.350	1.00 35.44
MOTA	3968	ND2	ASN B	121	47.917	-2.006	57.217	1.00 32.26
MOTA	3969	C	ASN B	121	51.172	-3.637	60.361	1.00 30.92
								1.00 27.08
ATOM	3970	0	ASN B		50.971		61.059	
atom	3971	N	VAL B	122	51.810	-2.560·	60.796	1.00 28.46
ATOM	3972	CA	VAL B	122	52.309	-2.457	62.155	1.00 29.48
ATOM	3973	CB	VAL B		53.840		62.177	1.00 33.40
ATOM	3974	CG1	VAL B		54.334	-2.294		1.00 32.22
ATOM	3975	CG2	VAL B		54.446		61.458	1.00 33.57
MOTA	3976	C	VAL B	122	51.713	-1.196	62.748	1.00 29.04
ATOM	3977	0	VAL B	122	51.800	-0.118	62.153	1.00 27.47
ATOM	3978	N	ALA, B		51.100	-1.326	63.918	1.00 26.71
MOTA	3979	CA	ALA B		50.477	-0.177	64.559	1.00 25.62
MOTA	3980	CB	ALA B		48.963	-0.281	64.447	1.00 21.65
MOTA	3981	С	ALA B	123	50.872	-0.005	66.017	1.00 28.62
ATOM	3982	ō	ALA B		51.227	-0.965	66.712	1.00 26.96
			_		50.805		66.472	1.00 22.85
ATOM	3983	N	PHE B	124	50.605	1.239	00.4/2	1.00 22.83

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- 500	3984	CA	PHE B	124		51.122	1.577	67.847	1.00 17.31
MOTA	3985	CB	PHE B			52.419	2.404	67.876	1.00 16.88
MOTA MOTA	3986	CG	PHE B			52.762	3.000	69.225	1.00 18.52
ATOM	3987		PHE B			52.533	2.304	70.403	1.00 17.52
ATOM	3988	CD2				53.382	4.245	69.297	1.00 17.88
ATOM-	3989	CE1	PHE B			52.914	2.837	71.638	1.00 25.77
ATOM	3990	CE2	PHE B			53.769	4.790	70.517	1.00 21.97
MOTA	3991	CZ	PHE B	124		53.535	4.084	71.698	1.00 20.16
ATOM	3992	c	PHE B			49.937	2.348	68.421	1.00 18.77
ATOM	3993		PHE B	124		49.462	3.311	67.820	1.00 16.62
ATOM	3994	N	ASN B	125		49.418	1.868	69.546	1.00 16.69
ATOM	3995	CA	ASN B	125		48.320	2.528	70.238	1.00 16.22
ATOM	3996	CB	ASN B	125	•	47.129	1.603	70.435	1.00 12.71
ATOM	3997	CG	ASN B			46.095	2.209	71.346	1.00 19.79 1.00 20.83
ATOM	3998	OD1				45.930	3.430	71.372 72.087	1.00 12.31
MOTA	3999		ASN B			45.376	1.371 3.004	71.600	1.00 19.19
MOTA	4000	С	ASN B			48.790 48.687	2.280	72.585	1.00 20.99
ATOM	4001	0	ASN B			49.335	4.226	71.668	1.00 19.02
MOTA	4002	N	PRO B		-	49.595	5.156	70.555	1.00 21.39
ATOM	4003	CD	PRO B			49.833	4.805	72.917	1.00 21.60
MOTA	4004	CA CB	PRO B			50.398	6.161	72.459	1.00 21.07
MOTA	4005	CG	PRO B			49.530	6.487	71.269	1.00 17.70
ATOM	4006 4007	C	PRO B			48.808	4.942	74.034	1.00 20.69
ATOM ATOM	4008	ō	PRO B			49.178	5.053	75.198	1.00 19.79
ATOM	4009	N	ALA B			47.525	4.937	73.689	1.00 16.67
ATOM	4010	CA	ALA B			46.476	5.065	74.698	1.00 20.44
	. 4011	CB	ALA B			45.198	5.609	74.066	1.00 19.56
ATOM	4012	С	ALA B	127		46.169	3.747	75.401	1.00 20.80
ATOM	4013	0	ALA B			45.555	3.742	76.472	1.00 19.47 1.00 20.52
MOTA	4014	N	GLY B			46.587	2.634	74.800 75.399	1.00 19.43
MOTA	4015	CA	GLY B			46.325	1.333 0.910	76.463	1.00 20.56
MOTA	4016	C	GLY B			47.327 48.182	1.697	76.869	1.00 18.37
ATOM	4017	0	GLY B			47.215	-0.333	76.929	1.00 19.68
ATOM	4018	N CA	GLY B			48.136	-0.820	77.943	1.00 19.93
MOTA	4019 4020	C	GLY B			47.620	-0.619	79.358	1.00 25.25
ATOM ATOM	4021	ō	GLY B			48.383	-0.686	80.329	1.00 18.98
ATOM	4022	N	MET B			46.317	-0.374	79.474	1.00 16.04
ATOM	4023	CA	MET B			45.677	-0.161	80.768	1.00 19.26
ATOM	4024	CB	MET B			44.301	0.451	80.519	1.00 17.94 1.00 22.95
MOTA	4025	CG	MET B			44.413	1.728	79.653 79.307	1.00 22.33
ATOM	4026	SD	MET B			42.873	2.615	78.382	1.00 20.22
ATOM	4027	CE	MET B			41.957	1.358 -1.548	81.421	1.00 22.63
ATOM	4028	C	MET B			45.598 44.546	-2.173	81.486	1.00 16.24
MOTA	4029	.0	MET B HIS B			46.737	-1.999	81.932	1.00 18.42
ATOM	4030	N CA	HIS B	131		46.853	-3.343	82.472	1.00 17.07
ATOM	4031 4032	CB	HIS B	131		48.323	-3.804	82.341	1.00 17.61
ATOM	4033	CG	HIS B			49.316	-2.979	83.106	1.00 14.01
atom Atom	4034		HIS B			49.138	-1.904	83.915	1.00 13.47
ATOM	4035	ND1	HIS B	131		50.680	-3.190	83.051	1.00 18.00
ATOM	4036	CE1	HIS B	131		51.297	-2.281	83.789	1.00 15.27
ATOM	4037	NE2	HIS B	131		50.384	-1.489	84.324	1.00 17.21 1.00 16.41
ATOM	4038	С	HIS B	131		46.329	-3.724	83.852	1.00 19.37
ATOM	4039	0	HIS B	131		46.452	-4.883	84.236 84.586	1.00 19.37
ATOM	4040	N	HIS B	132		45.721	-2.794	85.936	1.00 20.87
ATOM	4041	CA	HIS B			45.241	-3.112 -1.935	86.885	1.00 18.85
atom	4042	CB	HIS B			45.513	-1.686	87.152	1.00 20.00
ATOM	4043	CG	HIS B	132		46.966 47.715	-0.563	87.030	1.00 15.74
ATOM	4044	CDZ	HIS B	132		47.713	-2.655	87.659	1.00 14.72
ATOM	4045	WAT	HIS B	132		49.014	-2.139	87.837	1.00 14.64
ATOM	4046	MEG	HIS B	132		48.984	-0.872	87.462	1.00 14.88
ATOM	404.7 4048	C	HIS B	132		43.778	-3.547	86.136	1.00 22.83
arom	4049	ō	HIS B	132		43.478	-4.298	87.076	1.00 17.84
atom	7072	-							

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MOTA	4050	N	ALA E	3 133		42.878	-3.088	85.271	1.00 16.54
MOTA	4051	CA	ALA E	3 133		41.457	-3.396	85.424	1.00 19.13
ATOM	4052	CB	ALA E	3 133		40.654	-2.704	84.328	1.00 23.56
ATOM	4053	С	ALA E	133		41.127	~4.883	85.439	1.00 23.12
ATOM	4054	0	ALA E	133		41.718	-5.677	84.696	1.00 18.03
ATOM	4055	N	PHE E	134		40.181	-5.257	86.294	1.00 19.69
ATOM	4056	CA	PHE B			39.762	-6.649	86.365	1.00 19.35
MOTA	4057	CB	PHE B			39.583	-7.122	87.818	1.00 21.26
ATOM	4058	CG	PHE B			40.837	-7.053	88.646	1.00 23.41
ATOM	4059	CD1	PHE B			41.041	-6.009	89.544	1.00 24.25
ATOM	4060	CD2	PHE B			41.820	-8.027	38.522	1.00 22.80
MOTA	4061	CE1	PHE B			42.207	-5.935	90.311	1.00 23.36
ATOM	4062	CE2	PHE B			42.997	-7.964	89.283	1.00 27.74
ATOM	4063	cz	PHE B			43.190	-6.917	90.178	1.00 24.05
ATOM	4064	c	PHE B			38.444	-6.816	85.621	1.00 18.60
ATOM	4065	ŏ	PHE B			37.815	-5.849	85.196	1.00 13.82
	4066	N	LYS B			38.050	-8.064	85.454	1.00 19.78
MOTA	4067	CA	LYS B			36.813	-8.421	84.782	1.00 28.09
MOTA	4068		LYS B			36.501	-9.879	85.125	1.00 34.06
MOTA	4069		LYS B				-10.310	84.953	1.00 42.76
ATOM	4070		LYS B				-11.745	85.437	1.00 48.44
ATOM	4071		LYS B				-12.152	85.531	1.00 55.66
MOTA		•	LYS B				-11.332	86.544	1.00 51.65
MOTA	4072					35.639	-7.512	85.172	1.00 31.63
ATOM	4073		LYS B					84.309	1.00 24.86
MOTA	4074		LYS B			34.927	-6.999 -7.292	86.470	
MOTA	4075		SER B			35.450 34.331	-7.292 -6.477	86.933	1.00 29.89 1.00 30.86
MOTA	4076		SER B			33.282	-7.388	87.582	1.00 30.86
ATOM	4077		SER B					86.698	
MOTA	4078		SER B			32.916	-8.434	87.923	1.00 45.10 1.00 31.50
ATOM	4079		SER B			34.705	-5.380	88.765	
ATOM	4080		SER B			33.887	-4.997		1.00 24.54
MOTA	4081		ARG B			35.920	-4.854	87.835	1.00 22.63
ATOM	4082		ARG B			36.291	-3.826	88.794	1.00 25.51
ATOM	4083		ARG B			36.629	-4:486	90.136	1.00 29.62
ATOM	4084					36.391	-3.578	91.318	1.00 36.21
ATOM	4085		ARG B			86.874	-4.160	92.631	1.00 40.79
MOTA	4086					36.365	-3.357	93.744	1.00 45.95
MOTA	4087		ARG B			36.863	-3.369	94.973	1.00 41.97
MOTA	4088		ARG B			37.897	-4.144	95.263	1.00 43.42
ATOM	4089		ARG B			6.322	-2.604	95.913	1.00 46.65
MOTA	4090		ARG B			37.461	-2.956	88.339	1.00 24.73
MOTA	4091		ARG B			8.420	-3.441	87.734	1.00 19.32
MOTA	4092		ALA B			37.372	-1.663	88.631	1.00 16.77
MOTA	4093		ALA B			8.428	-0.733	88.270	1.00 18.50
ATOM	4094	-	ALA B			7.939	0.694	88.401	1.00 17.24
ATOM	4095		ALA B			9.597	-0.964	89.216	1.00 22.62
MOTA	4096		ALA B			9 411	-1.419	90.346	1.00 18.98
MOTA	4097		ASN B			0.301	-0.641	88.759	1.00 20.82
ATOM	4098		ASN B			1.989	-0.828	89.585	1.00 25.17
ATOM	4099		ASN B			2.311	-2.329	89.689	1.00 20.59
MOTA	4100		ASN B			3.556	-2.608	90.511	1.00 27.70
MOTA	4101		ASN B			3.726		91.592	1.00 22.43
MOTA	4102		ASN B			4.420	-3.487	90.010	1.00 24.43
MOTA	4103		ASN B			3.176	-0.062	89.020	1.00 22.37
ATOM	4104		ASN B			3.338	0.038	87.799	1.00 17.50
ATOM	4105		GLY B			3.984	0.496	89.920	1.00 21.67
ATOM	4106		GLY B			5.166	1.249	89.524	1.00 23.06
ATOM	4107		GLY B			5.005	2.268	88.402	1.00 26.29
ATOM	4108		GLY B			5.827	2.301	87.479	1.00 22.47
MOTA	4109		PHE B		4	3.958	3.093	88.473	1.00 22.33
ATOM	4110	CA F	PHE B	141	4	3.694	4.126	87.461	1.00 19.01
ATOM	4111	CB F	HE B	141	4	4.996	4.806	86.997	1.00 22.90
ATOM	4112	CG F	HE B	141	4	5.810	5.433	88.097	1.00 23.17
ATOM ·	4113	CD1 F	HE B	141	4	7.114	5.851	87.842	1.00 22.17
ATOM	4114	CD2 P	HE B	141	4	5.281	5.635	89.366	1.00 23.40
ATOM	4115	CE1 P	HE B	141		7.876	6.462	88.833	1.00 24.02
AION		- •							

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ATOM	4116	CE2 PHE B 141	46.033	6.244	90.361	1.00 23.03
	4117	CZ PHE B 141		6.658	90.092	1.00 25.15
ATOM					86.214	1.00 23.69
MOTA	4118	C PHE B 141		3.538		
MOTA	4119	O PHE B 141	42.596	4.283	85.335	1.00 18.88
MOTA	4120	N CYS B 142	42.962	2.211	86.122	1.00 15.03
				1.578	84.938	1.00 19.55
MOTA	4121					
ATOM	4122	CB CYS B 142	43.193	0.336	84.552	1.00 20.38
MOTA	4123	SG CYS B 142	44.933	0.662	84.190	1.00 37.40
	4124	C CYS B 142		1.171	85.098	1.00 22.77
MOTA				0.514	86.082	1.00 23.04
ATOM	4125	O CYS B 142				
ATOM	4126	N TYR B 143	40.094	1.557	84.130	1.00 15.24
MOTA	4127	CA TYR B 143	38.675	1.194	84.155	1.00 21.97
		CB TYR B 143		2.372	83.723	1.00 18.06
MOTA	4128					
MOTA	4129	CG TYR B 143		3.622	84.535	1.00 24.34
MOTA	4130	CD1 TYR B 143	39.038	4.516	84.214	1.00 23.20
ATOM	4131	CE1 TYR B 143	39.265	5.658	84.991	1.00 27.42
		CD2 TYR B 143	37.226	3.892	85.652	1.00 19.15
MOTA	4132					1.00 21.92
MOTA	4133	CE2 TYR B 143		5.023	86.432	
ATOM	4134	CZ TYR B 143	38.458	5.900	86.099	1.00 23.94
	4135	OH TYR B 143	38.655	7.015	86.877	1.00 22.37
MOTA				0.008	83.218	1.00 19.91
ATOM	4136					
MOTA	4137	O TYR B 143		-0.902	83.535	1.00 22.50
ATOM	4138	N ILE B 144	39.083	0.026	82.061	1.00 19.20
ATOM	4139	CA ILE B 144	38.938	-1.055	81.082	1.00 19.68
			38.282	-0.528	79.787	1.00 20.26
MOTA	4140					
MOTA	4141	CG2 ILE B 144		-1.649	78.760	1.00 15.37
ATOM	4142	CG1 ILE B 144	36.901	0.053	80.113	1.00 20.93
ATOM	4143	CD1 ILE B 144	36.198	0.697	78.917	1.00 23.75
			40.320	-1.627	80.774	1.00 22.78
MOTA	4144					1.00 22.01
MOTA	4145	O ILE B 144	41.281	-0.873	80.600	
ATOM	4146	N ASN B 145	40.422	-2.956	80.723	1.00 23.18
ATOM	4147	CA ASN B 145	41.698	-3.623	80.451	1.00 20.63
		CB ASN B 145	41.778	-4.935	81.243	1.00 17.81
MOTA	4148				81.268	1.00 25.17
MOTA	4149	CG ASN B 145	43.188	-5.531		
MOTA	4150	OD1 ASN B 145	43.804	-5.742	80.227	1.00 23.63
ATOM	4151	ND2 ASN B 145	43.693	-5.819	82.472	1.00 22.69
		C ASN B 145	41.780	-3.918	78.955	1.00 21.18
MOTA	4152					1.00 17.80
ATOM	4153	O ASN B 145	41.389	-5.002	78.508	
MOTA	4154	N ASN B 146	42.293	-2.968	78.177	1.00 15.23
ATOM	4155	CA ASN B 146	42.367	-3.175	76.733	1.00 19.71
		CB ASN B 146		-1.880	76.015	1.00 17.65
ATOM	4156				76.306	1.00 19.86
ATOM	4157	CG ASN B 146		-1.458		
ATOM	4158	OD1 ASN B 146	45.109	-1.735	75.532	1.00 20.27
ATOM	4159	ND2 ASN B 146	44.395	-0.798	77.435	1.00 11.85
	4160	C ASN B 146	43.277	-4.342	76.331	1.00 19.07
MOTA					75.328	1.00 18.61
MOTA .	4161	O ASN B 146	43.030	-4.996		
ATOM	4162	N PRO B 147	44.358	-4.598	77.082	1.00 17.78
ATOM	4163	CD PRO B 147	44.953	-3.919	78.240	1.00 18.13
	4164	CA PRO B 147	45.197	-5.735	76.678	1.00 19.98
ATOM				-5.694	77.698	1.00 24.29
MOTA	4165	CB PRO B 147	46.338			
ATOM	4166	CG PRO B 147	46.425	-4.201	78.020	1.00 26.27
ATOM	4167	C PRO B 147	44.377	-7.041	76.757	1.00 20.91
		O PRO B 147	44.461	-7.892	75.871	1.00 17.58
MOTA	4168				77.809	1.00 15.81
ATOM	4169	N ALA B 148	43.568	-7.172		
MOTA	4170	CA ALA B 148	42.732	-8.362	78.008	1.00 19.82
ATOM	4171	CB ALA B 148	42.049	-8,312	79.372	1.00 17.50
			41.683	-8.473	76.903	1.00 22.58
ATOM	4172				76.404	1.00 18.38
ATOM	4173	O ALA B 148	41.419	-9.567		
ATOM	4174	N VAL B 149	41.080	-7.341	76.540	1.00 22.48
	4175	CA VAL B 149	40.086	-7.300	75.466	1.00 19.04
ATOM			39.503	-5.877	75.281	1.00 18.96
ATOM	4176				73.988	1.00 17.32
MOTA	4177	CG1 VAL B 149	38.691	-5.800		1.00 17.32
ATOM	4178	CG2 VAL B 149	38.621	~5.531	76.462	1.00 15.33
	4179	C VAL B 149	40.763	-7.709	74.166	1.00 22.12
ATOM			40.240	-8.535	73.421	1.00 21.83
ATOM	4180				73.903	1.00 19.51
ATOM	4181	N GLY B 150	41.927	-7.120	, , , , , ,	T.00 T3.7T

ATOM 4183 C GLY B 150	ATOM	4182	CA	GLY B	150		42.657	-7.433	72.689	1.00 19.32
ATOM 4185 N ILE B 151 43.556 -9.435 73.700 1.00 19.51 ATOM 4186 CA ILE B 151 43.956 -10.834 73.723 1.00 23.25 ATOM 4187 CB ILE B 151 44.666 -11.175 75.053 1.00 23.50 ATOM 4188 CG2 ILE B 151 44.918 -12.679 75.158 1.00 21.98 ATOM 4190 CD1 ILE B 151 45.988 -10.394 75.129 1.00 21.98 ATOM 4191 C ILE B 151 45.988 -10.394 75.129 1.00 21.98 ATOM 4192 O ILE B 151 42.749 -11.741 73.490 1.00 28.24 ATOM 4193 N GLU B 152 40.647 -11.741 73.490 1.00 22.96 ATOM 4193 N GLU B 152 41.623 -11.450 74.144 1.00 27.32 ATOM 4195 CB GLU B 152 39.294 -11.845 74.886 1.00 26.46 ATOM 4196 CG GLU B 152 39.533 -12.200 76.347 1.00 28.26 ATOM 4196 CG GLU B 152 39.668 -14.181 75.617 1.00 28.26 ATOM 4198 OEI GLU B 152 39.668 -14.481 75.617 1.00 30.51 ATOM 4198 OEI GLU B 152 39.668 -14.481 75.617 1.00 30.51 ATOM 4190 OEZ GLU B 152 39.668 -14.481 75.617 1.00 30.53 ATOM 4200 C GLU B 152 39.948 -12.125 72.497 1.00 30.30 ATOM 4201 O GLU B 152 39.948 -13.042 77.767 1.00 30.53 ATOM 4202 N TYR B 153 39.948 -13.062 71.893 1.00 25.58 ATOM 4202 N TYR B 153 39.948 -13.062 71.893 1.00 25.62 ATOM 4203 CB TYR B 153 39.948 -13.062 71.893 1.00 25.62 ATOM 4204 CB TYR B 153 39.948 -13.022 71.893 1.00 25.63 ATOM 4204 CB TYR B 153 39.948 -13.062 71.893 1.00 25.69 ATOM 4204 CB TYR B 153 39.948 -13.062 71.893 1.00 25.69 ATOM 4204 CB TYR B 153 39.948 -13.062 71.893 1.00 25.69 ATOM 4205 CG TYR B 153 39.948 -13.062 71.893 1.00 25.69 ATOM 4204 CB TYR B 153 39.948 -13.062 71.893 1.00 25.69 ATOM 4204 CB TYR B 153 39.948 -13.062 71.893 1.00 25.69 ATOM 4205 CG TYR B 153 39.948 -13.062 71.893 1.00 25.69 ATOM 4204 CB TYR B 153 39.948 -13.066 68.200 1.00 27.94 ATOM 4204 CB TYR B 153 39.948 -13.062 71.893 1.00 25.69 ATOM 4205 CG TYR B 153 39.946 -13.062 71.893 1.00 25.69 ATOM 4207 CEI TYR B 153 39.946 -13.062 71.893 1.00 25.69 ATOM 4207 CEI TYR B 153 39.946 -13.062 71.94 71								-8.901	72.606	
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ATOM 4245 CG LYS B 157 40.157 -12.426 65.359 1.00 31.52 ATOM 4246 CD LYS B 157 39.132 -12.424 64.239 1.00 28.48							40.888	-13.746	65.418	1.00 29.79
ATOM 4246 CD LYS B 157 39.132 -12.424 64.239 1.00 28.48			CG	LYS B	157					
ATOM 4247 CE LYS B 157 38.295 -11.101 04.171 1.00 31.90	ATOM									
	ATOM	4247	CE	LYS B	15/		30.395	-11.101	. 04.1/1	1.00 31.90

181/263 Figure 18-65

- mov	4248	NZ	LYS B 157	37.406 -11.980	63.054	1.00 32.91
ATOM					65.381	
MOTA	4249		LYS B 157	40.724 -16.234		1.00 31.92
atom	4250		LYS B 157	41.146 -16.421	64.246	1.00 33.58
MOTA	4251		GLY B 158	40.890 -17.111	66.368	1.00 28.97
ATOM	4252	CA	GLY B 158	41.546 ~18.379	66.112	1.00 28.98
ATOM	4253	С	GLY B 158	42.962 -18.569	66.622	1.00 33.33
ATOM	4254		GLY B 158	43.503 -19.672	66.522	1.00 30.58
ATOM	4255		PHE B 159	43.578 -17.521	67.164	1.00 32.80
	4256		PHE B. 159	44.937 -17.657	67.678	1.00 28.89
ATOM						
ATOM	4257		- PHE B 159	45.560 -16.286	67.934	1.00 30.33
MOTA	4258		PHE B 159	45.748 -15.470	66.692	1.00 28:53
ATOM	4259		1 PHE B 159	44.682 ~14.787	66.121	1.00 24.58
ATOM	4260	CD:		46.989 -15.420	66.068	1.00 24.21
ATOM	4261	CE:	1 PHE B 159	44.849 -14.066	64.948	1.00 25.26
ATOM	4262	CE	PHE B 159	47.168 -14.706	64.895	1.00 23.66
ATOM	4263	CZ	PHE B 159	46.095 -14.026	64.332	1.00 26.65
ATOM	4264	С	PHE B 159	44.969 -18.484	68.958	1.00 30.92
ATOM	4265	ō	PHE B 159	44.102 -18.334	69.820	1.00 24.26
	4266	N	LYS B 160	45.979 -19.347	69.077	1.00 28.86
ATOM				46.123 -20.224	70.237	1.00 30.27
ATOM	4267	CA	LYS B 160			
MOTA	4268	CB	LYS B 160	46.085 -21.692	69.800	1.00 32.05
ATOM	4269	CG	LYS B 160	44.806 -22.117	69.113	1.00 41.13
ATOM	4270	CD	LYS B 160	. 44.809 -23.621	68.826	1.00 40.73
ATOM	4271	CE	LYS B 160	45.945 -24.031	67.904	1.00 43.16
ATOM	4272	NZ	LYS B 160	45.812 -23.408	66.554	1.00 48.69
ATOM	4273	С	LYS B 160	47.394 -19.997	71.048	1.00 28.23
ATOM	4274	0	LYS B 160	47.552 -20.561	72.130	1.00 25.29
ATOM	4275	N	ARG B 161	48.320 -19.206	70.520	1.00 28.51
ATOM	4276	CA	ARG B 161	49.550 -18.921	71.247	1.00 25.84
ATOM	4277	CB	ARG B 161	50.724 -19.719	70.667	1.00 25.33
ATOM	4278	CG	ARG B 161	50.551 -21.245	70.781	1.00 27.47
ATOM	4279	CD	ARG B 161	51.833 -21.985	70.394	1.00 32.27
	4280	NE	ARG B 161	52.218 -21.761	69.002	1.00 34.90
ATOM		CZ	ARG B 161	51.584 -22:276	67.954	1.00 38.45
ATOM	4281			50.527 -23.056	68.130	1.00 38.77
MOTA	4282	NH1		51.999 -22.000	66.725	1.00 38.64
ATOM	4283	NH2				
HOTA	4284	C	ARG B 161	49.818 -17.421	71.182	1.00 30.40
ATOM	4285	0	ARG B 161	50.393 -16.912	70.218	1.00 27.50
atom	4286	N	ILE B 162	49.376 -16.722	72.221	1.00 25.64
MOTA	4287	CA	ILE B 162	49.515 -15.273	72.303	1.00 27.44
MOTA	4288	CB	ILE B 162	48.13414.618	72.545	1.00 24.53
ATOM	4289	CG2		48.249 -13.101	72.473	1.00 25.49
ATOM	4290	CG1		47.142 -15.101	71.487	1.00 29.46
ATOM	4291	CD1		45.688 -14.707	71.758	1.00 31.94
ATOM	4292	С	ILE B 162	50. 465 -14.868	73.429	1.00 22.68
ATOM	4293	0	ILE B 162	5^.3 11 -15.3 0 2	74.568	1.00 24.25
ATOM	4294	N	LEU B 163	5454 -14.042	73.100	1.00 19.49
ATOM	4295	CA	LEU B 163	51.425 -13.561	74.081	1.00 17.57
ATOM	4296	CB	LEU B 163	53.8 50 - 13.686	73.528	1.00 20.54
ATOM	4297	CG	LEU B 163	54.979 -12.975	74.295	1.00 18.84
ATOM	4298		LEU B 163	55.102 -13.538	75.690	1.00 20.55
ATOM	4299		LEU B 163	56.293 -13.148	73.556	1.00 18.57
ATOM	4300	C	LEU B 163	52.158 -12.099	74.430	1.00 16.20
	4301	õ	LEU B 163	51.898 -11.277	73.549	1.00 16.09
ATOM			TYR B 164	52.227 -11.780	75.715	1.00 14.05
ATOM	4302	N		52.027 -10.411	76.191	
ATOM	4303	CA	TYR B 164			1.00 16.21
ATOM	4304	CB	TYR B 164	50.777 -10.323	77.070	1.00 16.01
ATOM	4305	CG	TYR B 164	50.534 -8.948	77.667	1.00 15.08
ATOM	4306	CD1		50.148 -7.869	76.869	1.00 19.51
ATOM	4307	CE1	TYR 5 164	49.948 -6.597	77.418	1.00 12.88
ATOM	4308	CD2	TYR B 164	50.715 -8.724	79.021	1.00 14.07
A.TOM	4309	CE2	TYR B 164	50.520 -7.463	79.583	1.00 13.66
FTOM	4310	CZ	TYR B 164	50.139 -6.407	78.782	1.00 14.72
ATOM	4311	он	TYR B .164	49.952 -5.163	79.354	1.00 13.54
ATOM	4312	С	TYR B 164	53.246 -10.017	77.018	1.00 19.14
ATOM	4313	ō	TYR B 164	53.539 -10.642	78.036	1.00 26.51
F-1-01;					-	

								76 577	1 00 22 40
MOTA	4314	N	ILE	В	165	53.964	-8.992	76.573	1.00 22.40
MOTA	4315	CA	ILE	В	165	55.148	-8.518	77.285	1.00 17.72
ATOM	4316	CB.	ILE	B	165	56.352	-8.465	76.343	1.00 22.51
		CG2	ILE		165	57.582	-7.902	77.079	1.00 16.36
MOTA	4317						-9.880	75.818	1.00 19.82
MOTA	4318	CG1	ILE		165	56.632			
ATOM	4319	CD1	ILE	В	165	57.721	-9.942	74.742	1.00 21.74
ATOM	4320	C	ILE	В	165	54.851	-7.126	77.850	1.00 22.54
ATOM	4321	0	ILE	В	165	54.478	-6.223	77.111	1.00 16.60
	4322	N	ASP		166	55.046	-6.961	79.156	1.00 15.78
ATOM			ASP		166	54.740	-5.704	79.840	1.00 20.62
MOTA	4323	CA			166	53.719	-5.996	80.949	1.00 17.57
MOTA	4324	CB	ASP				-4.742	81.486	1.00 25.39
MOTA	4325	CG	ASP		166	53.063			1.00 19.68
ATOM	4326		ASP		166	53.779	-3.859	82.003	
ATOM	4327	OD2	ASP	В	166	51.824	-4.637	81.377	1.00 29.22
ATOM	4328	C	ASP		166	55.976	-5.002	80.423	1.00 19.01
ATOM	4329	0	ASP		166	56.509	-5.412	81.456	1.00 19.74
	4330	N	LEU		167	56.414	-3.923	79.775	1.00 17.88
ATOM			LEU		167	57.598	-3.211	80.235	1.00 14.99
ATOM	4331	CA				58.412	-2.710	79.044	1.00 19.22
MOTA	4332	СВ	LEU.		167		-3.799	78.069	1.00 22.68
ATOM	4333	CG	LEU		167	58.871			1.00 25.35
ATOM	4334	CD1	LEU	В	167	59.835	-3.179	77.074	
ATOM	4335	CD2	LEU	В	167	59.570	-4.943	78.808	1.00 17.54
ATOM	4336	C ·	LEU	В	167	57.284	-2.059	81.183	1.00 17.49
ATOM	4337	0	LEU		167	58.189	-1.359	81.639	1.00 13.39
	4338	N	ASP		168	56.003	-1.878	81.479	1.00 20.03
ATOM		CA	ASP		168	55.549	-0.848	82.412	1.00 21.98
MOTA	4339				168	54.030	-0.955	82.597	1.00 21.21
MOTA	4340	CB	ASP				0.186	83.428	1.00 24.92
ATOM	4341	CG	ASP		168	53.453		83.753	1.00 22.98
MOTA	4342	С	ASP		168	56.241	-1.139		
ATOM	4343	0	ASP	В	168	56.447	-2.304	84.091	
ATOM	4344	OD1	ASP	В	168	52.849	1.099	82.825	1.00 22.03
ATOM	4345	OD2	ASP	В	168	53.606	0.189	84.676	1.00 18.43
ATOM	4346	N	ALA		169	56.581	-0.095	84.514	1.00 15.46
	4347	CA	ALA		169 -	57.263	-0.268	85.807	1.00 18.73
ATOM			ALA		169	57.764	1.084	86.323	1.00 11.98
ATOM	4348	CB			169	56.400	-0.940	86.886	1.00 21.82
ATOM	4349	C	ALA			56.886	-1.262	87.980	1.00 22.51
MOTA	4350	0	ALA		169			86.600	1.00 18.75
ATOM	4351	N	HIS		170	55.120	-1.134		1.00 22.70
MOTA	4352	CA	HIS		170	54.238	-1.776	87.570	
MOTA	4353	С	HIS	В	170	53.716	-3.096	87.015	1.00 22.11
ATOM	4354	0	HIS	В	170	53.536	-3.244	85.809	1.00 21.94
MOTA	4355	CB	HIS	В	170	53.050	-0.867	87.927	1.00 21.28
	4356	CG	HIS		170	53.449	0.475	88.460	1.00 18.89
ATOM	4357		HIS			53.695	1.539	87.626	1.00 19.13
MOTA			HIS			54.046	2.539	88.412	1.00 19.41
MOTA	4358					53.660	0.854	89.746	1.00 19.02
MOTA	4359	CD2	HIS	R	170			89.710	1,00 20.45
ATOM	4360	NE2	HIS	В	170	54.042	2.174		1.00 19.20
ATOM	4361	N	HIS	В	171	53.474	-4.047	87.907	
ATOM	4362	CA	HIS	В	171	52.961	-5.352	87.519	1.00 21.20
MOTA	4363	CB	HIS	В	171	52.964	-6.284	88.722	1.00 22.00
ATOM	4364	CG	HTS			52.541	-7.683	88.400	1.00 24.64
	4365		HIS			53.056	-8.594	87.540	1.00 19.19
ATOM		VD2	HIS	9	171	51.441	-8.279	88.979	1.00 25.71
ATOM	4366					51.295	-9.497	88.487	1.00 25.30
MOTA	4367		HIS					87.612	1.00 24.71
ATOM	4368	NE2	HIS	В	171	52.261	-9.713		1.00 23.91
MOTA	4369	С	HIS			51.549	-5.306	86.943	
ATOM	4370	0	HIS			50.677	-4.620	87.479	1:00 18.93
ATOM	4371	N	CYS.			51.332	-6.062	85.865	1.00 15.36
ATOM	4372	CA	CYS	3	172	50.036	-6.141	85.207	1.00 20.03
		CB	CYS	P	172	50.240	-6.534	83.732	1.00 22.46
ATOM	4373		CYS	P	172	51.259	-8.030	83.419	1.00 23.49
ATOM	4374	SG	CIS	D	172	49.110	-7.146	85.913	1.00 18.05
ATOM	4375	C	CYS	5	172			85.327	1.00 18.23
ATOM	4376	0	CYS	R	1/4	48.712	-8.151		1.00 16.78
ATOM	4377	N	ASP	В	173	48.767	-6.871	87.170	
ATOM	4378	CA	ASP	В	173	47.909	-7.776	87.928	
ATCM	1379	CB	ASP	В	173·	47.638	-7.236	89.344	1.00 20.39
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#### 183/263 Figure 18-67

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ASP B 173
                                    46.961
46.564
        4380
                                             -5.871
                                                               1.00 23.40
MOTA
               CG
                                                      89.354
                                             -5.435
MOTA
        4381
               OD1
                   ASP B 173
                                                      90.455
                                                               1.00 18.64
        4382
               OD2
                   ASP B 173
                                    46.834
                                             -5.231
                                                               1.00 19.24
MOTA
                                                      88.291
                                    46.595
MOTA
        4383
               C
                   ASP B 173
                                             -8.116
                                                      87.219
                                                               1.00 17.46
                   ASP B 173
                                    46.162
MOTA
        4384
               0
                                            -9.272
                                                      87.224
                                                               1.00 15.53
                   GLY B 174
                                    45.978
                                            -7.130
MOTA
        4385
               N
                                                      86.580
                                                               1.00 13.46
        4386
               CA
                   GLY B 174
                                    44.733
                                            -7.391
                                                      85.876
                                                               1.00 18.18
ATOM
                   GLY B 174
        4387
               С
                                    44.904
                                            -8.392
                                                      84.741
                                                               1.00 17.85
MOTA
                   GLY B 174
MOTA
        4388
               0
                                    44.104
                                            -9.316
                                                      84.583
                                                               1.00 18.27
ATOM
        4389
               N
                   VAL B 175
                                    45.951
                                            -8.214
                                                      83.943
                                                               1.00 16.14
        4390
                   VAL B 175
                                                               1.00 17.00
               CA
                                    46.206
                                            -9.111
                                                      82.829
MOTA
MOTA
        4391
               CB
                   VAL B 175
                                    47.305
                                            ~8.552
                                                      81.902
                                                               1.00 27.22
                   VAL B 175
                                    47.533
                                            ~9.507
        4392
               CG1
                                                      80.731
                                                               1.00 19.75
MOTA
        4393
              CG2
                   VAL B 175
                                    46.896
                                            -7.169
                                                      81.396
                                                               1.00 18.66
MOTA
        4394
               С
                   VAL B 175
                                    46.635 -10.486
                                                      83.324
                                                               1.00 22.82
MOTA
                   VAL B 175
                                    46.255 -11.503
        4395
               0
                                                      82.754
                                                               1.00 18.06
MOTA
        4396
                   GLN B 176.
                                                               1.00 21.67
ATOM
              N
                                    47.439 -10.520
                                                      84.378
ATOM
        4397
              CA
                   GLN B 176
                                    47.889 -11.798
                                                     84.911
                                                               1.00 21.55
                                    48.824 -11.602
        4398
              CB
                   GLN B 176
                                                      86.105
                                                               1.00 19.68
ATOM
                                    49.088 -12.905
50.066 -12.759
                                                     86.862
        4399
                                                               1.00 20.17
                   GLN B 176
MOTA
              CG
MOTA
        4400
              CD
                   GLN B 176
                                                     87.996
                                                               1.00 25.42
        4401
                   GLN B 176
                                    51.243 -12.442
                                                      87.786
                                                               1.00 21.56
              OE1
MOTA
        4402
                                                               1.00 20.18
              NE<sub>2</sub>
                   GLN B 176
                                    49.592 -13.000
                                                     89.217
ATOM
                   GLN B 176
                                    46.689 -12.630
                                                     85.348
ATOM
        4403
              C
                                                               1.00 24.78
                                                               1.00 22.91
ATOM
        4404
                   GLN B 176
                                    46.618 -13.817
                                                     85.057
                                   45.751 -12.007
44.571 -12.727
                   GLU B 177
        4405
              N
                                                     86.051
                                                              1.00 23.69
ATOM
        4406
              CA
                   GLU B 177
                                                     86.523
                                                              1.00 27.01
MOTA
                                    43.703 -11.825
        4407
              CB
                   GLU B 177
                                                     87.394
                                                              1.00 24.73
MOTA
                   GLU B 177
                                                                   37.46
ATOM
        4408
              CG
                                   42.633 -12.581
                                                     88.138
                                                              1.00
                                   41.767 -11.676
              CD
                   GLU B 177
                                                     88.987
                                                              1.00 42.48
MOTA
        4409
                                                             1.00 44.35
                  GLU B 177
                                                     88.432
ATOM
        4410
              OE1
                                   40.875 -11.002
                                   41.993 -11.627
ATOM
        4411
              OE2
                   GLU B 177
                                                     90.213
                                                               1.00
                                                                   45.63
        4412
              C
                   GLU B 177
                                   43.732 -13.247.
                                                     85.370
                                                              1.00 26.56
ATOM
                                                     85.408
                                                              1.00 27.71
                   GLU B 177
                                   43.240 -14.375
ATOM
        4413
              0
        4414
              N
                   ALA B 178
                                   43.573 -12.418
                                                     84.344
                                                              1.00 24.58
ATOM
                  ALA B 178
                                   42.776 -12.775
                                                     83.174
                                                              1.00 25.86
        4415
              CA
ATOM
              CB
                  ALA B 178
                                   42.778 -11.628
                                                     82.171
                                                              1.00 24.20
        4416
MOTA
                   ALA B 178
                                                     82.485
                                                              1.00
                                                                   25.72
ATOM
        4417
              C
                                   43.231 -14.054
        4418
                   ALA B 178
                                   42.406 -14.838
                                                     82.036
                                                              1.00 22.38
MOTA
              0
        4419
              N
                   PHE B 179
                                   44.535 -14.282
                                                     82.395
                                                              1.00 27.19
MOTA
                  PHE B 179
                                   44.990 -15.489
                                                     81.703
                                                              1.00 27.05
        4420
              CA
ATOM
ATOM
        4421
              CB
                  PHE B 179
                                   45.714 -15.086
                                                     80.418
                                                              1.00 25.22
MOTA
       4422
              CG
                  PHE B 179
                                   44.992 -14.020
                                                     79.644
                                                              1.00 20.36
                                   45.387 -12.687
                  PHE B 179
                                                     79.735
                                                              1.00 25.23
       4423
              CD1
ATOM
                                                     78.902
                  PHE B 179
                                                              1.00 19.22
ATOM
       4424
              CD2
                                   43.860 -14.332
ATOM
       4425
              CEL
                  PHE B 179
                                   44.659 -11.677
                                                     79.102
                                                              1.00 19.25
                                   43.128 -13.315
                                                     78.272
                  PHE B 179
                                                              1.00 20.65
       4426
              CE2
ATOM
                                                     78.374
                                                              1.00 25.64
                  PHE B 179
ATOM
       4427
              CZ
                                   43.528 -12.001
       4428
                  PHE B 179
                                   45.866 -16.398
                                                     82.556
                                                              1.00 23.50
MOTA
              С
                                                     82.038
                  PHE B 179
                                   46.652 -17.182
                                                              1.00 18.26
       4429
              0
ATOM
                                                     83.868
                  TYR B 180
                                                              1.00 23.24
ATOM
       4430
             N
                                   45.689 -16.313
                  TYR B 180
                                   46.479 -17.106
                                                     84.799
                                                              1.00
                                                                   26.76
       4431
             CA
MOTA
                                                     86.231
                  TYR B 180
                                   46.150 -16.665
                                                              1.00 25.72
             CB
       4432
MOTA
                                                     87.247
                                                              1.00 29.66
                  TYR B 180
                                   47.226 -16.969
MOTA
       4433
             CG
             CD1
                  TYR B 180
                                   47.037 -17.942
                                                     88.237
                                                              1.00 27.07
       4434
ATOM
                  TYR B 180
                                   48.039 -18.222
                                                     89.170
                                                              1.00 30.08
             CE1
       4435
ATOM
                                                     87.216
                                                              1.00 29.68
       4436
             CD2
                  TYR B 180
                                   48.444 -16.283
ATOM
                                   49.451 -16.552
49.248 -17.521
                  TYR B 180
                                                     88.139
                                                              1.00 30.99
       4437
             CE2
ATCM
                                                     89.112
                  TYR B 180
                                                              1.00 33.16
       4438
             CZ
MOTA
                                                     90.006
                                                              1.00 28.47
       1439
             OH
                  TYR B 180
                                   50.262 -17.791
MOTA
                  TYR B
                        180
                                   46.256 -18.619
                                                     84.649
                                                              1.00 29.13
       4440
             С
MOTA
                                                     84.922
                  TYR B 180
                                                              1.00 23.43
                                   47.163 -19.416
MOTA
       4441
             0
                                                     84.190
             N
                  ASP B 181
                                   45.073 -19.021
                                                              1.00 25.67
       4442
ATOM
                      В
                        181
                                                     84.075
                                                              1.00 28.28
       4443
             CA
                  ASP
                                   44.784 -20.445
MOTA
                                                     84.757
                  ASP B 181
                                   43.446 -20.759
                                                              1.00 32.13
       1444
             CB
ATOM
                                                     83.890
             CG
                  ASP B 181
                                   42.247 -20.410
                                                              1.00 36.12
       4445
ATCM
```

#### 184/263 Figure 18-68

						•
MOTA	4446	OD1 ASP B 181	42.202	-19.300	83.329	1.00 41.04
ATOM	4447	7 OD2 ASP B 181	41.334	-21.249	83.782	
MOTA	4448	3 C ASP B 181	44.773	-21.018	82.664	
MOTA	4449	O ASP B 181	44.246	-22.115	82.444	
ATOM	4450	N THR B 182		-20.302	81.702	1.00 29.24
ATOM	4451	CA THR B 182		-20.823	80.340	1.00 30.57
MOTA	4452			-20.008	79.397	1.00 30.03
MOTA	4453	OG1 THR B 182		-20.598	78.095	1.00 28.22
ATOM	4454	CG2 THR B 182		-18.561	79.310	1.00 26.55
ATOM	4455			-20.870	79.740	1.00 32.31
ATOM	4456	O THR B 182		-20.007	80.008	1.00 27.27
ATOM	4457			-21.878	78.909	1.00 29.94
MOTA	4458	CA ASP B 183		-22.049	78.273	1.00 31.40
ATOM	4459			-23.536	78.228	1.00 33.36
ATOM	4460	CG ASP B 183	47.718	-24.319	77.328	1.00 33.33
ATOM	4461	OD1 ASP B 183	46.513	-23.988	77.28 <del>7</del>	1.00 28.06
ATOM	4462	OD2 ASP B 183	48.186	-25.271	76.675	1.00 38.19
ATOM	4463	C ASP B 183	48.321	-21.462	76.864	1.00 31.14
MOTA	4464	O ASP B 183	49.332	-21.557	76.168	1.00 28.74
MOTA	4465	N GLN B 184		-20.852	76.446	1.00 25.34
ATOM	4466	CA GLN B 184	47.151	-20.251	75.118	1.00 28.59
MOTA	4467	CB GLN B 184	45.712	-20.256	74.581	1.00 26.84
ATOM	4468	CG GLN B 184	45.060	-21.632	74.529	1.00 34.86
ATOM	4469	CD GLN B 184	43.760	-21.647	73.736	1.00 32.27
ATOM	4470	OE1 GLN B 184	42.897	-20.789	73.912	1.00 35.43
ATOM	4471	NE2 GLN B 184	43.611	-22.641	72.870	1.00 28.92
ATOM	4472	C GLN B 184	47.672		75.175	1.00 27.28
MOTA	4473	O GLN B 184	47.871		74.148	1.00 29.70
ATOM	4474	N VAL B 185	47.900		76.386	1.00 27.64
ATOM	4475	CA VAL B 185	48.400		76.575	1.00 26.26
ATOM	4476	CB VAL·B 185	47.304		77.145	1.00 22.85
ATOM	4477	CG1 VAL B 185	47.879		77.395	1.00 23.10
ATOM	4478	CG2 VAL B 185	46.136		76.191	1.00 21.67
ATOM	4479	C VAL B 185	49.570		77.547	1.00 27.01
MOTA	4480 4481	O VAL B 185 N PHE B 186	49.456		78.663	1.00 23.75
ATOM	4482	N PHE B 186 CA PHE B 186	50.696		77.115	1.00 22.02
ATOM ATOM	4483	CB PHE B 186	51.868 53.142		77.978	1.00 21.83
ATOM	4484	CG PHE B 186	54.336		77.252 78.170	1.00 17.02 1.00 24.84
ATOM	4485	CD1 PHE B 186	54.756		78.580	1.00 22.70
ATOM	4486	CD2 PHE B 186	55.004		78.670	1.00 20.26
ATOM	4487	CE1 PHE B 186	55.819		79.471	1.00 20.20
ATOM	4488	CE2 PHE B 186	56.071		79.563	1.00 20.01
ATOM	4489	CZ PHE B 186	56.481		79.968	1.00 17.84
ATOM	4490	C PHE B 186	52.032		78.368	1.00 18.12
ATO:	4491	O PHE B 186	52.038		77.508	1.00 15.92
ATO.4	4492	N VAL B 187	52.161 -		79.661	1.00 18.06
ATOm	4493	CA VAL B 187	52.348 -		80.153	1.00 17.67
ATOM	4494	CB VAL B 187	51.282 -		81.225	1.00 22.85
MOTA	4495	CG1 VAL B 187	51.608 -	-11.473	81.840	1.00 24.08
ATOM	4496	CG2 VAL B 187	49.882 -	-12.808	80.598	
MOTA	4497	C VAL B 187	53.735 -		80.788	1.00 18.32
ATOM	4498	O VAL B 187	54.092 -	-13.807	81.707	1.00 18.82
ATOM	4499	N LEU B 188	54.503 -	-12.103·	80.282	1.00 14.70
MOTA	4500	CA LEU B 188	55.832 -	11.789	80.798	1.00 18.84
ATCM	4501	CB LEU B 188	56.900 -	11.948	79.716	1.00 18.64
ATOM	450,2	CG LEU B 188	58.230 -	11.277	80.082	1.00 21.23
ATOM	4503	CD1 LEU B 188	58.769 -	11.832	81.395	1.00 18.55
MOTA	4504	CD2 LEU B 188	59.227 ~	11.489	78.957	1.00 20.49
ATOM	4505	C LEU B 188	55.836 -	10.339	81.280	1.00 22.14
ATOM	4506	O LEU B 188		-9.410	80.517	1.00 19.96
ATOM	4507	N SER B 189		10.133	82.540	1.00 21.08
ATOM	4508	CA SER B 189	56.203	-8.782	83.961	1.00 21.85
ATOM	4509	CB SER B 189		-8.543	83.908	1.00 25.95
ATCM	4510	OG SER B 189		-7.252	84.475	1.00 21.91
ATOM	4511	C SER B 189	57.423	-8.420	83.883	1.00 23.62

**185/263** Figure 18-69

* mov	4512	0	.SER E	100		57 020	0 174	04 256	1 00 10 61
ATOM						57.829	-9.174	84.766	1.00 18.61
ATOM	4513		LEU E			58.020	-7.269	83.569	1.00 20.83
ATOM	4514	CA	LEU E	190		59.149	-6.767	84.347	1.00 21.85
ATOM	4515	CB	LEU E	190		60.278	-6.226	83.473	1.00 22.85
	4516		LEU B			60.964	-7.089	82.413	
ATOM									1.00 32.59
MOTA	4517					62.337	-6.479	82.140	1.00 29.27
MOTA	4518	CD2	LEU B	190		61.136	-8.511	82.979	1.00 31.88
ATOM	4519	С	LEU B	190		58.505	-5.613	85.085	1.00 21.28
	4520		LEU B						
_ATOM						57.695	-4.897	84.501	1.00 15.72
ATOM	4521	N	- HIS B	191		58.857	-5.421	86.351	1.00 18.16
MOTA	4522	CA	HIS B	191		58.249	-4.357	87.145	1.00 17:46
ATOM	4523	CB	HIS B	191		56.759	-4.690	87.369	1.00 16.00
-	4524	CG	HIS B			56.517			
ATOM							-6.085	87.880	1.00 22.14
ATOM	4525	CD2				56.341	-6.551	89.143	1.00 12.25
ATOM	4526	ND1	HIS B	191		56.372	-7.179	87.049	1.00 18.02
MOTA	4527	CE1	HIS B	191		56.119	-8.256	87.775	1.00 8.17
	4528	NE2				56.094	-7.902	89.049	1.00 19.79
ATOM									
ATOM	4529	С	HIS B			58.945	-4.197	88.484	1.00 17.41
ATOM	4530	0	HIS B	191		59.769	-5.029	88.867	1.00 18.74
'ATOM	4531	N	GLN B	192		58.618	-3.114	89.182	1.00 18.20
ATOM	4532	CA				59.173	-2.854	90.502	1.00 18.41
ATOM	4533	CB				58.690	-1.500	91.034	1.00 20.71
ATOM	4534	CG	GLN B	192		58.871	-0.334	90.072	1.00 21.49
ATOM	4535	CD	GLN B	192		58.226	0.930	90.594	1.00 20.65
ATOM	4536	OE1	GLN B	192		58.775	1.615	91.459	1.00 21.52
	4537	NE2	-	192		57.029	1.226	90.098	1.00 15.10
ATOM									
ATOM	4538	С		192		58.608	-3.945	91.395	1.00 17.55
ATOM	4539	0	GLN B	192	•	57.415	-4.256	91.320	1.00 17.48
ATOM	4540	N	SER B	193		59.447	-4.522	92.240	1.00 15.71
ATOM	4541	CA	SER B			58.986	-5.574	93.143	1.00 20.58
					•		-5.963		
MOTA	4542	CB		193		60.093		94.120	1.00 20.71
MOTA	4543	OG	SER B			59.571	-6.804	95.138	1.00 22.55
ATOM	4544	С	SER B	193		57.774	-5.112	93.947	1.00 21.81
ATOM ·	4545	0	SER B	193		57.769	-4:003	94.486	1.00 20.82
ATOM	4546	N	PRO B			56.745	-5.967	94.063	1.00 21.80
								•	
ATOM	4547	CD		194		56.648	-7.331	93.524	1.00 24.27
ATOM	4548	CA		194		55.524	-5.643	94.812	1.00 23.58
MOTA	4549	CB	PRO B	194		54:678	-6.909	94.642	1.00 22.98
ATOM	4550	CG	PRO B	194		55.168	-7.458	93.317	1.00 26.35
ATOM	4551	С		194		55.841	-5.366	96.283	1.00 25.79
-									
ATOM	4552	9		194		55.009	-4.831	97.022	1.00 27.26
MOTA	4553	N	GLU B	195		57.045	-5.736	96.710	1.00 23.20
ATOM	4554	CA	GLU B	195		57.428	-5.514	98.093	1.00 29.56
ATOM	4555	CB	GLU B	195		58.816	-6.090	98.379	1.00 32.38
	4556	CG		195		58.940	-7.567		1.00 45.25
ATOM									
MOTA	4557	CD		195		60.206	-8.189	98.613	1.00 50.44
MOTA	4558	OE1	GLU B	195		61.290	-7.580	98.471	1.00 50.51
MOTA	4559	OE2	GLU B	195		60.118	-9.297	99.184	1.00 49.77
ATOM	4560	C	GLU B	195		57.414	-4.035	98.425	1.00 25.11
	4561		GLU B			57.095	-3.659	99.551	1.00 29.05
ATOM			-	-					
ATOM	4562		TYR B			57.729		97.445	1.00 22.90
ATOM	4563	CA	TYR B	196 .		57.743	-1.750	97.696	1.00 22.46
ATCM	4564	CB	TYR B	196		59.188	-1.223	97.668	1.00 22.72
	.4565		TYR B			59.855	-1.234	96.301	1.00 24.17
ATOM	4566		TYR B			59.639			
ATOM							-0.203	95.385	1.00 20.87
MOTA.	4567		TYR B	196		60.229	-0.222	94.118	1.00 18.31
ATOM	4568	CD2	TYR B	196		60.684	-2.289	95.916	1.00 24.63
ATOM	4569		TYR B			61.276	-2.318	94.648	1.00 24.39
	4570		TYR B			61.042	-1.284	93.756	1.00 23.01
ATOM									
atom	4571		TYR B :			61.592	-1.328	92.492	1.00 19.86
ATOM	4572	C '	TYR B	196		56.896	-0.938	96.725	1.00 23.54
ATOM	4573		TYR B	196		56.779	0.275	96.869	1.00 17.53
	4574		ALA B			56.293	-1.589	95.740	1.00 22.11
ATOM									
ATOM	4575		ALA B			55.503	-0.829	94.779	1.00 24.28
ATOM	4576		ALA B			56.310	-0.616	93.513	1.00 23.03
ATOM	4577	C 2	ALA B 1	197		54.153	-1.412	94.413	1.00 22.80
								-	

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#### Figure 18-70

ATOM	4578	3 0	ALA B	197	53.91	0 -2.609	94.549	1 00	17.67
ATOM	4579		PHE B	198	53.27		93.932		26.40
ATOM	4580	) CA	PHE B	198	51.95		93.495		28.19
ATOM	4581	. CB		198	51.15		93.035		29.51
ATOM	4582	. CG	PHE B	198	49.72		92.711		29,50
MOTA	4583	CD	1 PHE B	198	48.73		93.674		32.77
ATOM	4584	CD i	2 PHE B	198	49.36	7 -0.533	91.455		25.82
ATOM	4585	CE	1 PHE B	198	47.41	0 -0.223	93.394		36.70
ATOM	4586	CE	2 PHE B.	198	48.05	0 -0.858	91.170		29.29
MOTA	4587		PHE B	198	47.07		92.141	1.00	33.05
MOTA	4588		PHE B		52.17	0 -1.858	92.284		28.28
ATOM	4589			198	53.04		91.456	1.00	27.15
MOTA.	4590			199	51.40		92.185		31.37
MOTA	4591		PRO B		51.44		91.045		37.07
MOTA	4592		PRO B		50.38		93.144		35.32
MOTA	4593		PRO B		49.54		92.321		33.88
MOTA	4594	CG	PRO B		50.64		91.578		36.75
MOTA	4595 4596	C O	PRO B	199	51.24		94.184		36.93
MOTA MOTA	4597	N	PHE B		52.30		93.860		50.93
MOTA	4598	CA	PHE B		50.804 51.644		95.422 96.421		37.04
ATOM	4599			200	51.547		97.723		30.13
MOTA	4600	CG		200	51.760		97.543		28.70 29.98
ATOM	4601	CD1		200	50.717		97.137		28.92
ATOM	4602	CD2			53.016		97.746		23.60
ATOM	4603	CE1			50.922		96.938		27.63
MOTA	4604	CE2	PHE B 2	200	53.229		97.547		23.56
MOTA	4605	CZ	PHE B 2	200	52.182		97.143		28.37
MOTA	4606	С	PHE B 2	200	51.296	-6.227	96.658		25.51
MOTA	4607	0	PHE B 2		52.112	-6.984	97.167	1.00	20.92
ATOM	4608	N	GLU B 2		50.094		96.252		27.41
ATOM	4609	CA	GLU B 2		49.576		96.454		31.98
ATOM	4610	CB	GLU B 2		48.056		96.487		31.57
MOTA	4611	CG	GLU B 2		47.486		97.449		39.17
MOTA	4612 4613	CD OE1		201	45.987		97.316		40.31
ATOM_ ATOM	4614	OE2		01	45.332 45.475		97.500 97.019		38.90 35.04
ATOM	4615	C		01	49.979		95.422		30.83
ATOM	4616	ō		01		-10.219	95.690		26.34
MOTA	4617	N		02	50.362		94.234		24.95
ATOM	4618	CA		02	50.764		93.195		22.79
MOTA	4619	CB	LYS B 2	02	49.588		92.258		25.12
MOTA	4620	CG	LYS B 2	02	48.484	-10.523	93.000		35.38
MOTA	4621	CD		02		-11.099	92.103	1.00	38.67
MOTA	4622	CE	LYS B 2			-11.998	92.903		40.98
ATOM	4623	NZ	LYS B 2			-12.659	92.028		46.65
MOTA	4624	C	LYS B 2		51.975		92.435	1.00	
MOTA	4625	0	LYS B 2		52.355		92.549		21.83
MOTA	4626	N	GLY B 2		52.598	-9.910	91.684	1.00	
ATOM	4627 4628	CA C	GLY B 2		53.779	-9.545 -10.297	90.928 91.396	1.00	
MOTA MOTA	4629	o	GLY B 2			-10.297	90.888	1.00	
ATOM	4630	Ŋ	PHE B 2	04		-11.201	92.358	1.00	
ATOM	4631	CA	PHE B 2			-11.957	92.859	1.00	
ATOM	4632	CB	PHE B 2			-12.567	94.236	1.00	
ATOM	4633	CG	PHE B 20			-11.549	95.322	1.00	
ATOM	4634	CD1	PHE B 20			-10.977	95.535	1.00	
ATOM	4635	CD2	PHE B 20			-11.159	96.133	1.00	
ATOM	4636	CEl	PHE B 20			-10.036	96.545	1.00	
MOTA	4637	CE2	PHE B 20	04		-10.221	97.141	1.00	
ATOM	<b>463</b> €	CC	PHE B 20		55.124	-9.658	97.350	1.00	25.54
ATOM	4639	C	PHE B 20		56.412	-13.057	91.894	1.00 2	
ATCM	4640	0	PHE B 20		55.613	-13.540	91.091	1.00	
MOTA	4641		LEU B 20		57.676	-13.449	91.986	1.00	
MOTA	4642		LEU B 20		58.233	-14.472	91.114	1.00	
ATOM	4643		LEU B 20	15	59.723	-14.637	91.413	1.00	94.UI
			U 1						

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#### Figure 18-71

MOTA	4644	CG LEU 3 205	60.495 -15.669	90.592	1.00 34.12
ATOM	4645	CD1 LEU 3 205	60.356 -15.382	89.109	1.00 32.95
MOTA	4646		61.957 -15.629	91.005	1.00 36.49
MOTA	4647		57.535 -15.827	91.205	1.00 30.51
ATOM	4648		57.467 -16.562	90.220	1.00 25.89
ATOM	4649		57.010 -16.147	92.382	1.00 30.43
ATOM	4650		56.338 -17.423	92.605	1.00 30.64
MOTA	4651	CB GLU B 206	56.025 -17.601	94.093	1.00 34.77
MOTA	4652 4653	CG GLU B 206 CD GLU B 206	57.227 -17.512	95.033	1.00 42.50
ATOM ATOM	4654	OE1 GLU B 206	57.718 -16.084 58.228 -15.438	95.270	1.00 45.76
ATOM	4655	OE2 GLU B 206	57.585 -15.602	94.333	1.00 42.62 1.00 50.22
ATOM	4656	C GLU B 206	55.045 -17.587	91.811	1.00 30.22
ATOM	4657	O GLU B 206	54.607 -18.708	91.563	1.00 28.18
MOTA	4658	N GLU B 207	54.430 -16.472	91.425	1.00 25.16
MOTA	4659	CA GLU B 207	53.178 -16.499	90.664	1.00 28.78
MOTA	4660	CB GLU B 207	52.546 -15.107	90.695	1.00 30.76
ATOM	4661	CG GLU B 207	52.121 -14.659	92.093	1.00 29.39
ATOM	4662	CD GLU B 207	52.057 -13.151	92.230	1.00 27.87
ATOM	4663	OE1 GLU B 207	51.656 -12.477	91.261	1.00 24.38
ATOM	4664	OE2 GLU B 207	52.389 -12.636	93.316	1.00 25.36
MOTA	4665	C GLU B 207	53.453 -16.922	89.224	1.00 29.48
MOTA MOTA	4666 4667	O GLU B 207 N ILE B 208	53.658 -16.077	88.351	1.00 27.48
ATOM	4668	CA ILE B 208	53.442 -18.230 53.735 -18.754	88.976 87.646	1.00 26.67 1.00 32.60
ATOM	4669	CB ILE B 208	54.789 -19.877	87.740	1.00 32.80
ATOM	4670	CG2 ILE B 208	55.239 -20.296	86.352	1.00 41.65
ATOM	4671	CG1 ILE B 208	56.008 -19.404	88.532	1.00 36.07
ATOM	4672	CD1 ILE B 208	56.814 ~18.338	87.851	1.00 45.18
MOTA	4673	C ILE B 208	52.522 -19.289	86.870	1.00 32.26
MOTA	4674	O ILE B 208	52.668 -19.799	85.759	1.00 27.43
ATOM	4675	N GLY 5 209	51.328 -19.165	87.442	1.00 32.60
ATOM	4676	CA GLY B 209 C GLY B 209	50.139 ~19.652	86.760	1.00 35.07
ATOM ATOM	4677 4678	C GLY B 209 . O GLY B 209	49.565 -20.892 50.230 -21.524	87.420 88.235	1.00 36.19
ATOM	4679	N GLU B 210	48.335 -21.245	87.066	1.00 31.61
ATOM	4680	CA GLU B 210	47.677 -22.412	87.647	1.00 40.60
ATOM	4681	CB GLU B 210	46.633 -21.964	88.672	1.00 37.98
ATOM	4682	CG GLU B 210	45.446 -21.234	88.058	1.00 42.78
ATOM	4683	CD GLU B 210	44.470 -20.717	89.098	1.00 48.41
ATOM	4684	OE1 GLU B 210	43.400 -20.202	88.709	1.00 51.03
MOTA	4685	OE2 GLU B 210	44.778 -20.814	90.306	1.00 49.90
ATOM	4686	C GLU B 210	46.996 -23.248	86.564	1.00 39.48
ATOM ATOM	4687 4688	O GLU B 210 N GLY B 211	46.709 -22.751 46.736 -24.515	85.471 86.876	1.00 33.65
ATOM	4689	CA GLY B 211	46.087 -25.399	85.923	1.00 39.18 1.00 38.43
ATOM	4690	C GLY B 211	46,877 -25.500	84.637	1.00 40.29
ATOM	4691	O GLY B 211	48.101 -25.610	84.666	1.00 39.39
ATOM	4692	N LYS B 212	46.187 -25.458	83.504	1.00 40.90
MOTA	4693	CA LYS E 212	46.864 -25.538	82.219	1.00 43.53
MOTA	4694	CB LYS B 212	45.842 -25.548		1.00 47.87
ATOM	4695	CG LYS B 212	44.795 -26.665	81.144	1.00 53.09
ATOM	4696	CD LYS B 212	45.398 -28.076	81.130	1.00 58.61
ATOM	4697	CE LYS B 212	46.069 -28.454	82.452	1.00 59.78
ATOM	4698	NZ LYS B 212 C LYS B 212	46.670 -29.825	82.420	1.00 62.17
ATOM ATOM	4699 4700	C LYS B 212 O LYS B 212	47.823 -24.363 48.797 -24.457	82.040 81.295	1.00 38.84
ATOM	4701	N GLY B 213	47.543 -23.262	82.731	1.00 37.20
ATOM	4702	CA GLY B 213	48.384 -22.081	82.627	1.00 37.20
ATOM	4703	C GLY B 213	49.625 -22.107	83.505	1.00 37.09
ATOM	4704	O GLY B 213	50.425 -21.165	83.489	1.00 25.85
ATOM	4705	N LYS B 214	49.794 -23.180	84.273	1.00 33.33
MOTA	4706	CA LYS B 214	50.953 -23.297	85.148	1.00 37.90
ATOM	1707	CB LYS B 214	50.886 -24.598	85.954	1.00 38.89
MOTA	4708	CG LYS B 214	52.032 -24.786	86.938	1.00 39.29
ATOM	4709	CD LYS B 214	51.876 -26.094	87.704	1.00 43.60

ATOM	4710	CE	LYS E	214	53 047	-26.334	88.640	1.00 47.36
				_				
$MOT \mathcal{K}$	4711	NZ	LYS E	214	23.102	-25.264	89.666	1.00 54.03
MOTA	4712	С	LYS E	214	52.209	-23.275	84.291	1.00 34.16
				•				
ATOM	4713	0	LYS E	214	52.404	-24.136	83.438	1.00 34.70
MOTA	4714	14	GLY B	215	53.057	-22.279	84.523	1.00 33.58
MOTA	4715	CA	GLY B	212	54.275	-22.152	83.743	1.00 28.27
MOTA	4716	С	GLY B	215	54.104	-21.155	82.605	1.00 31.02
			-					
ATOM	4717	0	GLY B	215	55.033	-20.911	81.833	1.00 23.68
MOTA	4718	N	TYR B	216	52 918	-20.564	82.493	1.00 22.45
MOTA	4719	CA	TYR B	216	52.683	-19.605	81.426	1.00 24.03
MOTA	4720	CB	TYR B	216 -	51 458	-20.013	80.603	1.00 17.60
MOTA	4721	CG	TYR B	216	51.682	-21.291	79.806	1.00 25.96
ATOM	4722	CD1	TYR B	216	51 692	-22.538	80.435	1.00 21.41
							-	
ATOM	4723	CE1	TYR B	216	51.988	-23.704	79.729	1.00 23.78
ATOM	4724	CD2	TYR B	216	51 970	-21,242	78.439	1.00 19.62
ATOM	4725	,CE2	TYR B	210	52.269	-22.402	77.72 <del>2</del>	1.00 26.39
ATOM	4726	CZ	TYR B	216	52 277	-23.630	78.379	1.00 29.35
ATOM	4727	ОН	TYR B	210	52.5//	-24.782	77.690	1.00 27.75
ATOM	4728	C -	TYR B	216	52.575	-18.153	81.884	1.00 24.53
ATOM	4729	0	TYR B	210	J2.U0J	-17.298	81.159	$1.00^{\circ}$ 18.99
ATOM	4730	N	ASN B	217	53.052	-17.886	83.098	1.00 21.41
ATOM	4731	ÇA	ASN B	21/	53.073	-16.534	83.642	1.00 21.23
ATOM	4732	CB	ASN B	217	51.954	-16.325	84.669	1.00 16.78
MOTA	4733	CG	ASN B		21.882	-14.889	85.162	1.00 22.07
ATOM	4734	OD1	ASN B	217	52,506	-14.521	86.163	1.00 23.13
ATOM	4735	MDZ	ASN B			-14.058	84.435	
ATOM	4736	С	ASN B	217	- 54 . 437	-16.339	84.291	1.00 19.40
	4737		ASN B	217	54 057	-17.145	85.124	1.00 19.28
ATOM		0						
ATOM	4738	N	LEU B	218	55.130	-15.273	83.905	1.00 18.65
	4739	CA	LEU B	219	56 450	-15.004	84.444	1.00 16.41
MOTA								
ATOM	4740	CB	LEU B	218	57.512	-15.244	83.368	1.00 18.29
ATOM	4741	CG	LEU B	218	58 851	-15.872	83.782	1.00 28.15
ATOM	4742	CDI	LEU B	718	59.873	-15.563 _.	82.695	1.00 20.50
ATOM	4743	CD2	LEU B	218 .	59 332	-15.348	85 116	1.00 22.53
MOTA	4744	С	LEU B	218	56.595	-13.562	84.926	1.00 17.89
ATOM	4745	0	LEU B	218	56.469	-12.627	84.128	1.00 14.48
ATOM	4746	N	ASN B	219		-13.395	86.219	1.00 14.09
ATOM	4747	CA	ASN B	219	57.044	-12.075	86.821	1.00 18.41
MOTA	4748	СВ	ASN B			-11.922	88.111	1.00 14.64
MOTA	4749	CG	ASN B	219	54.748	-11.898	87.868	1.00 27.12
			ASN B				86.880	1.00 20.21
MOTA	4750					-11.332		
ATOM	4751	ND2	ASN B	219	53.982	-12.480	88.787	1,00 23.62
	4752	С	ASN B	210		-11.843	87.172	1.00 20.39
ATOM	•							
MOTA	4753	0	ASN B	219	59.115	-12.672	87.841	1.00 20.41
MOTA	4754	N	ILE B	220	59 056	-10.717	86.729	1.00 15.11
MOTA	4755	CA	ILE B.	.22C	60.441	-10.394	87.033	1.00 17.16
ATOM	4756	CB	ILE B	22C	61.250	-10.083	85.740	1.00 20.78
MOTA	4757	CG2	ILE B		62.736	-9.821	86.094	1.00 18.08
MOTA	4758	CG1	ILE B	220	61.138	-11.250	84.748	1.00 17.62
							85.273	
ATOM	4759	CD1				-12.590		1.00 20.72
ATOM	4760	С	ILE B	220	60.475	-9.161	87.947	1.00 21.17
			ILE B		60.565	-8.036	87.470	.1.00 16.03
MOTA	4761	0						
MOTA	4762	N	PRO B	221	60.367	-9.357	89.274	1.00 21.74
		CD	PRO B			-10.619	90.000	1.00 22.96
ATOM	4763							
ATOM	4764	CA	PRO B	2 <b>21</b>	60.394	-8.225	90.213	1.00 19.16
		CB	PRO B		59.947	-8.869	91.523	1.00 19.40
ATOM	4765							
ATOM	4766	CG	PRO B	221	60.564	-10.251	91.407	1.00 23.02
		С	PRO B		61.799	-7.634	90.289	1.00 22.42
MOTA	4767							
MOTA	4768	0	PRO B	221	.62.780	-8.369	90.425	1.00 20.71
		N	LEU B	222	61.899	-6.309	90.202	1.00 22.74
ATOM	4769							
ATOM	4770	CA	LEU B	222	63.198	-5.643	90.223	1.00 21.18
	4771	CB	LEU B	222	63.453	-4.993	88.850	1.00 17.21
ATOM							87.721	
ATOM	4772	CG	LEU B	444	63.467	-6.027		1.00 20.26
	4773	CD1	LEU B	222	63.453	-5.354	86.361	1.00 20.00
ATOM		257		222			87.881	
ATCM :	4774	CD2	LEU B		64.696	-6.908		1.00 21.93
••-	4775	С	LEU B		63.335	-4.616	91.353	1.00 20.04
ATOM	4	_					•	-

### **189/263** Figure 18-73

```
ATOM
        4776
               0
                   LEU B 222
                                    62.350
                                             -4.030
                                                     91.806
                                                              1.00 17.58
                   PRO B 223
                                    64.571
                                             -4.394
                                                     91.830 . 1.00 19.48
ATOM
        4777
                                                     91.400
        4778
               CD
                   PRO B 223
                                    65.806
                                             ~5.072
MOTA
                                                               1.00 16.80
ATOM
        4779
               CA
                   PRO B 223
                                    64.873
                                             -3.454
                                                     92.915
                                                               1.00
                                                                    20.38
        4780
               CB
                   PRO B 223
                                    66.274
                                             -3.881
                                                     93.327
                                                               1.00 26.11
ATOM
        4781
               CG
                   PRO B 223
                                    66.884
                                             -4.161
                                                     91.973
MOTA
                                                               1.00 19.74
        4782
               C
                   PRO B 223
                                    64.818
                                             -1.971
                                                     92.553
MOTA
                                                               1.00 21.39
        4783
               0
                   PRO B 223
                                    64.815
                                             -1.598
                                                     91.380
                                                               1.00 17.16
ATOM
ATOM
        4784
               N
                   LYS B 224
                                    64.798
                                             -1.142
                                                     93.589
                                                               1.00 20.65
                   LYS B 224
                                    64.755
        4785
               CA.
                                             0.311
                                                     93.462
                                                               1.00 27.00
MOTA
                                    64.577
                                             0.938
               CB
                   LYS B 224
                                                     94.844
        4786
ATOM
                                                               1.00
                                                                    36:47
        4787
               CG
                   LYS .B 224
                                    63.415
                                              0.389
                                                     95..651
                                                               1.00 37.72
ATOM
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MOTA
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               CD
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                                              0.833
                                                               1.00 42.06
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ATOM
                   LYS B 224
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        4790
               NZ
                   LYS B 224
                                    62.645
                                              0.570
                                                     99.399
MOTA
                                                               1.00 46.30
        4791
               C
                   LYS B 224
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                                                     92.874
                                                               1.00 27.01
MOTA
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ATOM
        4792
              0
                   LYS B 224
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                                                     92.995
                                                               1.00 21.54
        4793
                   GLY B 225
                                    66.038
                                              1.989
                                                     92.259
                                                               1.00 22.66
MOTA
              N
                   GLY B 225
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                                             2.565
                                                     91.669
                                                               1.00 25.67
ATOM
        4794
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                                             1.809
MOTA
        4795
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                   GLY B 225
                                    67.768
                                                     90.459
                                                               1.00
                                                                    24.95
        4796
              Ö
                   GLY B 225
                                    68.917
                                             1.975
                                                     90.069
                                                               1.00 26.83
ATOM
                                                               1.00 21.79
        4797
                   LEU B 226
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                                             0.980
                                                     89.855
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                                                               1.00
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                   LEU B 226
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                                            -0.473
                                                     88.099
                                                               1.00 23.29
ATOM
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                                                     87.091
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        4802
              CD2:LEU B 226
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                                            -1.997
                                                     86.508
                                                               1.00 22.96
ATOM
                   LEU B 226
                                    68.008
                                             1.017
                                                     87.603
                                                               1.00
ATOM
        4803
              c
                                                                    22.66
                   LEU B 226
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                                             2.087
                                                     87.250
                                                               1.00 20.19
ATOM
        4804
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                   ASN B 227
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                                                     87,060
                                                               1.00 15.52
ATOM
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        4806
              CA
                  ASN B 227
                                    69.794
                                             1.317
                                                     85.998
                                                               1.00 19.49
ATOM
                  ASN B 227
                                    71.304
                                                     86.270
                                                               1.00 20.43
        4807
              CB
                                             1.474
ATOM
                  ASN B 227
                                   72.062
                                                     86.206
                                                               1.00 28.97
        4808
              CG
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ATOM
                                                               1.00 24.30
                                                     85.199
MOTA
        4809
              OD1
                  ASN B 227
                                    72.015
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                  ASN B 227
                                   72.786
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ATOM
        4810
MOTA
        4811
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                  ASN B 227
                                    69.548
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                                                     84.630
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                                   69.004
                                                              1.00 18.90
                                            -0.432
                                                     84.555
ATOM
        4812
              0
                  ASN B
                         227
                  ASP B 228
                                   69.949
                                             1.347
                                                     83.552
                                                              1.00 20.98
MOTA
        4813
              N
                                   69.720
                                                     82.208
                                                               1.00 22.61
        4814
              CA
                  ASP
                      3
                         228
                                             0.817
ATOM
                  ASP B 228
                                   70.270
                                             1.753
                                                     31.126
                                                               1.00 23.46
              CB
ATOM
        4815
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                                                               1.00 26.12
ATOM
        4816
              CG
                  ASP 3 228
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                                                     81.119
                  ASP B
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                                             3.193
                                                     81.415
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ATOM
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              OD1
        4818
              OD2
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                                             4.101
                                                     80.773
                                                               1.00 30.22
ATOM
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                                                               1.00 23.49
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                  ASP B 228
                                   70.286
ATOM
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              C
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                  ASP B
                         228
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                                                     81,288
                                                              1.00 19.31
ATOM
       4821
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                  ASN B 229
                                   71.484
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                                                     82.453
                                                               1.00 22.24
ATCM
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                                                              1.00 23.30
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ATOM
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                                                     82.737
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                                                              1.00 20.99
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ATOM
       4825
              OD1
                                                              1.00 27.44
ATOM
       4826
              ND2
                  ASN B
                         229
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                                                     82.417
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       4827
                  ASN B 229
                                   71.341
                                                     82,943
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ATCM
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MOTA
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       4829
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                  GLU B 230
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              CA
                  GLU B 230
                                   70.069
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ATOM
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              CB
                  GLU 3 230
ATOM
       4831
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MOTA
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              CG
                  GLU B
                         230
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                                                     88.470
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       4833
              CD
                  GLU B 230
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ATOM
                                                              1.00 27.87
                                                     88.569
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             OE1
                  GLU B 230
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ATOM
       4834
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       4835
              OE2
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ATOM
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                  GLU B 230
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ATOM
       1836
             С
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                                                     84.022
                  GLU B 230
                                   68.347
                                            -5.445
       4837
             О
ATOM
                                                              1.00 21.46
                        231
       :838
             N
                  PHE B
                                   68.091
                                            -3.242
                                                     83.637
ATCM
                                                     82.933
                                                              1.00 22.84
                  PHE B 231
                                   66.814
                                            -3.429
             CA
       4839
ATOM
                                                              1.00 23.96
                                            -2.079
                  PHE B 231
                                                     82.529
                                   66.210
       4840
             CB
atom
                  PHE B
                        231
                                   64.803
                                                     81.975
                                                              1.00 26.13
       4841
             CG
                                            -2.182
ATOM
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									4 00 05 00
ATOM	4842	CD1	.PHE B	231		63.738	-2.514	82.805	1.00 25.00
		CD2	PHE B			64.550	-1.956	80.627	1.00 22.93
ATOM	4843								
MOTA	4844	CE1	PHE B	231		62.440	-2.618	82.304	1.00 25.03
	4845	CE2	PHE B	231		63.250	-2.059	80.114	1.00 27.46
MOTA								80.957	1.00 20.25
ATOM	4846	cz	PHE B	231		62.196	-2.390		
MOTA	4847	С	PHE B	231		66.978	-4.288	81.677	1.00 23.14
			PHE B			66.221	-5.239	81.464	1.00 20.02
ATOM	4848	0							
MOTA	4849	N	LEU B	232		67.963	-3.952	80.845	1.00 22.02
		CA	LEU B			68.200	-4.697	79.614	1.00 19.97
MOTA	4850				•				
MOTA	4851	CB	LEU B	232		69.192	-3.942	78.734	1.00 24.99
ATOM	4852	CG	LEU B	232		68.665	-2.581	78.263	1.00 29.73
								77.454	1.00 28.11
ATOM	4853	CD1				69.746	-1.856		
MOTA.	4854	CD2	LEU B	232		67.409	-2.784	77.414	1.00 26.54
			LEU B			68.688	-6.119	79.898	1.00 19.25
MOTA	4855	С							
ATOM	4856	0	LEU B	232		68.365	-7.051	79.162	1.00 19.49
	4857	N	PHE B	233		69.468	-6.280	80.962	1.00 20.50
MOTA									
MOTA	4858	CA	PHE B	233		69.950	-7.599	81.378	1.00 20.70
MOTA	4859	CB	PHE B	233		70.825	-7.471	82.632	1.00 23.75
						71.217	-8.790	83.239	1.00 28.58
MOTA	4860	CG	PHE B						
ATOM	4861	CD1	PHE B	233		72.285	-9.519	82.731	1.00 30.48
ATOM	4862	CD2	PHE B	233		70.481	-9.328	84.294	1.00 25.32
	-							83.262	1.00 31.87
ATOM	4863	CE1	PHE B			72.617			
ATOM	4864	CE2	PHE B	233		70.803	-10.573	84.832	1.00 31.26
	4865	CZ	PHE B			71 871	-11.292	84.317	1.00 32.29
MOTA									1.00 20.23
MOTA	4866	C	PHE B	233	•	68.712	-8.439	81.727	
MOTA	4867	0	PHE B	233		68.553	-9.567	81.270	1.00 21.56
	4868	N	ALA B			67.842	-7.878	82.560	1.00 21.26
MOTA								82.963	1.00 19.60
MOTA	4869	CA	ALA B		٠.		-8.576		
ATOM	4870	CB	ALA B	234		65.835	-7 <i>.</i> 733	83.950	1.00 19.25
MOTA	4871	С	ALA B	234		65.772	-8.898	81.749	1.00 18.87
						65.253	-10.010	81.624	1.00 21.91
MOTA	4872	0	ALA B						
MOTA	4873	N	LEU B	235		65.634	-7.934	80.845	1.00 20.29
MOTA	4874	CA	LEU B	235		64.822	-8.141	79.652	1.00 19.53
		CB	LEU B			64.773	-6:874	78 [.] .795	1.00 24.07
MOTA	4875							78.024	1.00 27.87
MOTA	4876	CG	LEU B			63.465	-6.607		
ATOM	4877	CD1	LEU B	235		63.783	-5.813	76.770	1.00 20.77
	4878	CD2	LEU B	235		62.761	-7.897	77.664	1.00 26.94
MOTA							-9.276	78.795	1.00 20.79
MOTA	4879	С	LEU B			65.376			
MOTA	4880	0	LEU E	235		64.648	-10.205	78.431	1.00 18.25
ATOM	4881	N	GLU B	236		66.665	-9.191	78.462	1.00 19.33
					•	67.303	-10.206	77.629	1.00 27.93
ATOM	4882	CA	GLU B						
ATOM	4883	CB	GLU B	236		68.7 <b>7</b> 7	-9.853	77.384	1.00 31.06
ATOM	4884	CG	GLU B	236		68.969	-8.597	76.548	1.00 43.60
		CD	GLU B			70.428	-8.292	76.259	1.00 45.19
ATOM	4885								1.00 48.77
ATOM	4886	OE1	GLU B	236		70.697	-7.309	75.538	
ATOM	4887	OE2	GLU B	236		71.300	-9.032	76.751	1.00 52.72
			GLU B			67.20_	-11.607	78.209	1.00 24.89
ATOM	4888	C						77.501	1.00 22.06
ATOM	4889	0	GLU B	236		66.865	-12.552		
ATOM	4890	N	LYS B	237		67.520	-11.748	79.492	1.00 24.20
	4891	CA	LYS B				-13.058	80.130	1.00 27.10
MOTA								81.562	1.00 23.43
ATOM	4892	CB	LYS B	237		67.989	-12.984		
MOTA	4893	CG	LYS B	237		69.466	-12.641	81.650	1.00 29.46
	4894	CD	LYS B				-13.683	80.924	1.00 31.65
MOTA								80.993	1.00 39.70
MOTA	4895	CE	LYS B	237			-13.356		
MOTA	4896	NZ	LYS B	237		72.580	-14.363	80.242	1.00 46.74
	4897	C	LYS B			66.019	-13.615	80.143	1.00 30.92
MOTA								79.763	1.00 31.42
MOTA	4898	0	LYS B	237			-14.766		1.00 05 00
ATOM	4899	N	SER B	238		65.057	-12.806	80.573	1.00 25.86
		CA	SER B			63.677	-13.280	80.620	1.00 27.98
MOTA	4900							81.289	1.00 23.89
MOTA	4901	CB	SER B			62.776	-12.241		1.00 20.00
ATOM	4902	OG	SER B	238		62.756	-11.028	80.565	1.00 29.27
	1903		SER E	238		63.145	-13.642	79.229	1.00 28.32
MOTA		С						79.089	1.00 29.65
MOTA	4904	0	SER B	238			-14.605		1 00 27 30
ATOM	4905	N	LEU B	239		63.536	-12.886	78.203	1.00 27.39
	4906	CA	LEU B	239			-13.192	76.846	1.00 32.52
ATOM								75.837	1.00 30.53
MOTA	4907	CB	LEU B	233		03.344	-12.129		2.00 00.0-
AION									

ATOM	4908	CG LEU B 239	62, 833	-10.77	2 75.895	1.00 36.06
ATOM	4909	CD1 LEU B 239	63.404			
ATOM	4910	CD2 LEU B 239	61.338	_		1.00 30.73
MOTA	4911	C LEU B 239	63.598		3 76.430	1.00 34.15
MOTA	4912	O LEU B 239		-15.340		1.00 33.23
ATOM	4913 4914	N GLU B 240		~14.859		
MOTA ATOM	4915	CA GLU B 240 CB GLU B 240	65.434	-16.152	76.472	
ATOM	4916	CG GLU B. 240	67 070	-16.238 -15.407	77.011	
ATOM	4917	CD GLU B 240		-15.532		
ATOM	4918	OE1 GLU B 240		-16.679		1.00 48.20 1.00 45.36
MOTA	4919	OE2 GLU B 240		-14.492		1.00 48.21
ATOM	4920	C GLU B 240	64.604	-17.258	77.108	1.00 33.23
ATOM	4921	O GLU B 240		-18.310		1.00 32.15
ATOM	4922	N ILE B 241		-17.017		1.00 29.72
ATOM ATOM	4923 4924	CA ILE B 241 CB ILE B 241		-17.989		1.00 29.85
ATOM	4925	CG2 ILE B 241	63.006	-17.489 -18.456	80.466	1.00 30.42
ATOM	4926	CG1 ILE B 241	64 309	-17.311	81.162 81.254	1.00 29.88
ATOM	4927	CD1 ILE B 241		-16.760		1.00 29.77 1.00 32.92
ATOM	4928	C ILE B 241		-18.247	78.298	1.00 34.65
MOTA	4929	O ILE B 241	61.592	-19.396	78.149	1.00 30.22
MOTA		N VAL B 242	61.379	-17.178	77.823	1.00 31.16
MOTA MOTA		CA VAL B 242	60.114	-17.312	77.105	1.00 34.55
ATOM		CB VAL B 242 CG1 VAL B 242	59.476 58.191	-15.937	76.825	1.00 30.77
ATOM		CG2 VAL B 242	59.201		76.038 78.140	1.00 32.18
ATOM		C VAL B 242	60.320		75.787	1.00 31.57 1.00 36.56
MOTA		O VAL B 242		-18.959	75.453	1.00 33.93
ATOM		N LYS B 243	61.337		75.042	1.00 38.64
ATOM		CA LYS B 243	61.659	-18.241	73.760	1.00 44.36
ATOM ATOM		CB LYS B 243 CG LYS B 243	62.966	-17.659	73.214	1.00 48.33
ATOM		CD LYS B 243	62.810 62.185		72.386	1.00 53.88
ATOM		CE LYS B 243	63.056	-10.718 -17 681	71.036 70.242	1.00 53.72 1.00 54.69
ATOM	4943	NZ LYS B 243	62.456	-18.025	68.923	1.00 57.75
MOTA		LYS B 243	61.793	-19.755	73.824	1.00 43.21
MOTA		LYS B 243	61.432	-20.455	72.884	1.00 42.92
ATOM ATOM		N GLU B 244 CA GLU B 244	62.312 -		74.935	1.00 45.77
ATOM		GLU B 244	62.528 - 63.669 -		75.085	1.00 47.72
ATOM		G GLU B 244	64.080		76.075 76.208	1.00 50.89 1.00 57.16
ATOM		D GLU B 244	65.223 -		77.173	1.00 57.16
ATOM		E1 GLU B 244	66.295 -		76.942	1.00 60.18
ATOM		E2 GLU B 244	65.049 -	-24.308	78.160	1.00 61.59
ATOM ATOM	4953 C			-22.507	75.505	1.00 47.78
ATOM	4954 O			-23.736	75.544	1.00 51.39
ATOM		A VAL B 245	60.200 - 59.019 -	.22.831 .22.500	75.805 76.230	1.00 43.31
ATOM		B VAL B 245	58.867 -	22.514	77.771	1.00 43.55 1.00 45.89
ATOM	4958 C	G1 VAL B 245 .			78.231	1.00 49.90
ATOM		G2 VAL B 245	60.131 -	23.040	78.435	1.00 46.37
ATOM	4960 C		57.727 -	22.115	75.565	1.00 41.01
ATOM	4961 0		56.659 -	22.676	75.798	1.00 39.36
ATOM ATOM	4962 N 4963 C		57.814 -		74.716	1.00 34.37
ATOM	4964		56.610 - 55.986 -		74.077 74.958	1.00 34.36
MOTA	4965 C		54.542 ~		74.538	1.00 30.80 1.00 32.57
MOTA		D1 PHE B 246	53.548 -		74.989	1.00 32.37
ATOM		2 PHE B 246	54.174 -		74.003	1.00 28.20
ATOM		E1 PHE B 246	52.207 -	19.878	74.704	1.00 26.71
MOTA		E2 PHE B 246	52.836 -		73.713	1.00 29.27
ATOM	4970 CC	PHE B 246	51.850 -	18.689	74.065	1.00 26.18
MOTA MOTA	4971 C 4972 O	PHE B 246	56.904 -: 57.740 -:	20.040	72.682 72.517	1.00 35.32
	4973	GLU B 247	56.205 -	20 268 20 268	71.683	1.00 31.73
			50.205 -			1.00 37.00

### **192/263** Figure 18-76

ATOM				56.363	-20.137	70.296	1.00 40.73
MOTA				56.518	-21.347	69.370	
ATOM				56.670	-22.702	70.073	
ATOM				55.381	-23.214	70.718	1.00 55.29
MOTA				54.887	-22.601		
ATOM	4979			54.859	-24.246	70.241	
MOTA	4980			55.090	-19.379	69.939	
ATOM ATOM	4981 4982			54.129	-19.960	69.436	
ATOM	4983			55.076	-18.064 -17.270		
ATOM	4984		_	53 035	-17.188	70.733	
ATOM	4985		-	54 375	-17.188	69.916	
ATOM	4986			55.844	-15.880	70.562 70.233	1.00 35.33 1.00 32.85
ATOM	4987			53.563	-16.990	68.457	
MOTA	4988	O PRO B 248		54.427	-16.808	67.604	1.00 29.38
ATOM	4989	N GLU B 249		52.263	-17.012	68.182	1.00 32.23
ATOM	4990	CA GLU B 249		51.773	-16.782	66.828	1.00 29.35
ATOM	4991	CB GLU B 249		50.374	-17.366	66.645	1.00 31.87
ATOM	4992	CG GLU B 249		50.284	-18.867	66.787	1.00 28.64
ATOM	4993	CD GLU B 249	-	48.847	-19.338	66.747	1.00 33.37
ATOM	4994	OE1 GLU B 249		48.069	-18.917	67.630	1.00 26.38
ATOM	4995 4996	OE2 GLU B 249		48.494	-20.115	65.835	1.00 37.71
ATOM	4997	C GLU B 249 O GLU B 249		51.700	-15.273	66.650	1.00 28.25
ATOM	4998	N VAL B 250		51 561	-14.765 -14.564		1.00 21.47
ATOM	4999	CA VAL B 250		51.361	-13.110	67.768 67.756	1.00 21.77
ATOM	5000	CB VAL B 250		50.027	-12.676	67.357	1.00 21.41 1.00 26.42
MOTA	5001	CG1 VAL B 250		49.037	-13.196	68.378	1.00 20.96
ATOM	5002	CG2 VAL B 250		49.931	-11.166	67.243	1.00 24.68
ATOM	5003	C VAL B 250		51.757	-12.608	69.168	1.00 22.88
ATOM	5004	O VAL B 250		51.592	-13.354	70.133	1.00 18.00
ATOM	5005	N TYR B 251			-11.359	69.295	1.00 20.28
ATOM	5006	CA TYR B 251		52.481	-10.823	70.620	1.00 22.33
MOTA	5007 5008	CB TYR B 251 .		53.956	-11.043	70.999	1.00 20.67
ATOM ATOM	5009	CG TYR B 251 CD1 TYR B 251		54.948		70.427	1.00 21.89
ATOM	5010	CE1 TYR B 251		55.198 56.129	-8.834 -7.922	71. <b>07</b> 2 70.5 <b>6</b> 2	1.00 19.31
ATOM	5011	CD2 TYR B 251			-10.321	69.254	1.00 23.73 1.00 18.72
ATOM	5012	CE2 TYR B 251		56.580	-9.417	68.734	1.00 18.72
ATOM	5013	CZ TYR B 251		56.813	-8.220	69.390	1.00 27.33
ATOM	5014	OH TYR B 251		57.705	-7.308	68.865	1.00 23.18
ATOM	5015	C TYR B 251		52.134	-9.349	70.732	1.00 25.71
ATOM	5016	O TYR B 251		52.095	-8.622	69.728	1.00 20.14
MOTA	5017	N LEU B 252		51.834	-8.930	71.958	1.00 21.13
MOTA MOTA	5018 5019	CA LEU B 252		51.533	-7.532	72.252	1.00 24.61
ATOM	5020	CB LEU B 252 CG LEU B 252		50.154	-7.373	72.897	1.00 22.88
ATOM	5021	CD1 LEU B 252		48.915 48.779	-7.435 -8.792	1.996	1.00 23.73
ATOM	5022	CD2 LEU B 252		47.697	-7.119	72.833	1.00 23.18
ATOM	5023	C LEU B 252		52.610	-7.044	73.217	1.00 24.77
ATOM	5024	O LEU B 252				74.076	1.00 23.33
MOTA	5025	N LEU B 253		53.011	-5.786	73.071	1.00 20.14
MOTA		CA LEU B 253		54.057	-5.209	73.911	1.00 20.33
MOTA		CB LEU B 253		55.304	-4.946	73.051	1.00 15.18
MOTA		CG LEU B 253		56.490	-4.210	73.688	1.00 18.34
ATOM		CD1 LEU B 253		57.062	-5.044	74.829	1.00 14.11
MOTA	* .	CD2 LEU B 253		57.552	-3.953	72.624	1.00 19.60
MOTA		C LEU B 253		53.550	-3.913	74.536	1.00 20.54
ATOM ATOM		O LEU B 253 N GLN B 254		53.200	-2.974	73.821	1.00 22.80
ATOM		N GLN B 254 CA GLN B 254		53.495 53.000	-3.858	75.865 76.539	1.00 20.37 1.00 21.77
ATOM		CB GLN B 254		52.129	-2.654 -3.040	76.339 77.755	1.00 21.77
ATOM		CG GLN B 254			-3.040 -2.815	79.124	1.00 17.65
ATOM ·		CD GLN B 254			-1.396	79.609	1.00 28.19
ATOM		DE1 GLN B 254			-0.996	80.124	1.00 26.96
ATOM	5039	NE2 GLN B 254			-0.619	79.432	1.00 16.80
						•	

ATOM	5040	С	GLN	B 254		54.211	-1.793	76.887	1.00 20.15
	5041	0		B 254		55.186	2.254	77.497	1.00 20.11
ATOM			_						
ATOM	5042	N	. LEU	B 255		54.146	-0.532	76.468	1.00 19.46
ATOM	5043	CA	T ETT	в 255		55.268	0.386	76.614	1.00 15.99
							_		
ATOM	5044	CB	LEU	B 255		55.692	0.831	75.211	1.00 18.15
ATOM	5045	CG	LEH	B 255		56.143	-0.316	74.296	1.00 21.80
MCTA	5046	CDI	LEU	B 255		56.215	0.159	72.850	1.00 16.70
ATOM	5047	CD2	LEU	B 255		57.501	-0.843	74.771	1.00 13.76
								77.492	
MOTA	5048	С		B 255		55.083	1.614		1.00 21.41
ATCM	5049	0	_ LEU	B 255		55.379	2.741	77.065	1.00 18.40
	5050	N	GLY			54.618	1.408	78.718	1.00 16.80
ATOM									
ATOM	5051	CA	GLY	B 256		54.456	2.519	79.634	1.00 19.90
ATOM	5052	С	GT.Y	B 256	•	55.816	3.181	79.818	1.00 17.68
_									
ATOM	5053	0	GLY	B 256		56.854	2.514	79.841	1.00 13.96
MOTA	5054	N	THR	B 257		55.824	4.497	79.936	1.00 19.55
	5055	CA		B 257		57.081	5.205	80.098	1.00 19.47
MOTA						_			
ATOM	5056	CB		B 257		57.044	6.547	79.340	1.00 21.49
MOTA	5057	OC1	THR	B 257		55.989	7.365	79.858	1.00 17.43
ATOM	5058	CG2	THR	B 257		56.780	6.311	77.850	1.00 22.49
ATOM	5059	C	THR	B 257		57.440	5.466	81.564	1.00 20.75
	5060	0		B 257		58.480	5.054	81.843	1.00 25.01
MOTA									
MOTA	5061	N	ASP	B 258		56.618	5.004	82.504	1.00 17.23
MOTA	5062	CA	ASP	B 258		56.929	5.277	83.906	1.00 17.42
	5063	CB	ASP			55.744	4.940	84.846	1.00 12.75
ATOM									
ATOM	5064	CG	ASP			55.197	3.524	84.676	1.00 21.60
MOTA	5065	С	ASP	B 258		58.245	4.718	84.460	1.00 16.09
	5066	0		B 258		58.667	5.116	85.542	1.00 22.07
ATOM									
ATOM	5067		ASP :			55.901	2.642	84.150	1.00 17.74
ATOM	5068	OD2	ASP :	B 258		54.041	3.281	85.109	1.00 18.68
ATOM	5069	N	PRO	B 259		58.879	3.746	83.779	1.00 20.98
					•		2.901	82.641	1.00 17.75
ATOM	5070	CD		B 259		58.474	2.901		
ATOM	5071	CA	PRO :	B 259		60.154	3.257	84.321	1.00 22.63
MOTA	5072	CB	PRO :	B 259		60.395	1.988	83.506	1.00 23.46
		CG		B 259		59.800	2:343	82.199	1.00 27.08
MOTA	5073								
MOTA	5074	С	PRO	в 259		61.305	4.284	84.172	1.00 23.86
MOTA	5075	0	PRO 1	3 259		62.406	4.082	84.698	1.00 24.24
	5076	N		3 260		61.054	5.387	83.465	1.00 20.49
ATOM									
MOTA	5077	CA		B 260		62.080	6.417	83.262	1.00 15.17
ATOM	5078	CB	LEU I	3 260		61.626	7.408	82.185	1.00 17.03
ATOM	5079	CG	LETT 1	3 260		61.431	6.881	80.760	1.00 16.02
									1.00 17.03
ATOM	5080		LEU I			60.703	7.915	79.901	
ATOM	5081	CD2	LEU 1	3 260		62.803	6.546	80.163	1.00 18.58
ATOM	5082	С	LEU 1	3 260		62.449	7.194	84.541	1.00 22.45
						61.611	7.440	85.412	1.00 17.84
ATOM	5083	0		3 260					
atom	5084	N	LEU 1	3 261		63.713	7.588	84.635	1.00 22.90
ATOM	3085	CA	LEU I	3 261		64.219	8.332	85.782	1.00 26.34
		CB	LEU I			65.605	8.914	85.473	1.00 20.58
ATOM	5086								
ATOM	<i>5</i> 087	CG	LEU 1	3 261		66.180	9.850	86.553	1.00 28.44
ATOM	5088	CD1	LEU I	3 261		66.481	9.055	87.812	1.00 29.84
	5089		LEU I			67.462	10.522	86.057	1.00 32.10
atom		_							
ATOM	5090	С	LEU I	3 261			9.475		1.00 27.61
ATOM	5091	0	LEU I	3 261		62.978	9.586	87.408 .	1.00 24.02
	5092		GLU I			62.934	10.315	85.269	1.00 23.33
ATOM		Ŋ							
MOTA	5093	CA	GLU I	3 262		62.126	11.490	85.530	1.00 23.38
ATOM	5094	CB	GLU E	262		62.115	12.415	84.302	1.00 23.17
	5095	CG	GLU I			63.503	12.854	83.806	1.00 28.98
ATOM									
ATOM	5096	CD	GLU E			64.179	11.831	82.902	1.00 32.26
ATOM	5097	CE1	GLU I	262		63.702	10.673	82.838	1.00 29.28
	5098		GLU I			65.201	12.186	82.264	1.00 25.42
atom								85.976	
ATOM	5099	C	GLU E			60.693	11.249		1.00 23.25
ATOM	5100	0	GLU E	262		60.013	12.192	86.368	1.00 27.63
	5101	N	ASP E	263		60.219	10.011	85.927	1.00 22.25
ATOM		CA	ASP E			58.840	9.751	86.345	1.00 24.46
ATOM	5102					_			
ATOM	5103	CB	ASP F	263		58.214	8.659	85.465	1.00 20.94
ATOM	5104	CG	ASP E	263		56.710	8.543	85.659	1.00 25.30
			ASP E			55.995	8.318	84.656	1.00 21.82
ATOM	5105		-10F E	. 203			0.220	•	

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MOTA	5106	OD2 ASP B 26	56.239	8.666	86.811	1.00 18.31
ATOM	5107					
				9.339		1.00 26.39
MOTA	5108		· ·	8.335		1.00 22.11
MOTA	5109			10.124	88.648	1.00 25.81
ATOM	5110	CA TYR B 26	4 58.101	9.864	90.084	1.00 30.96
MOTA	5111	CB TYR B 26	4 57.511	11.055	90.841	1.00 36.80
MOTA	5112	CG TYR B 26	4 58.241	12.356		1.00 46.58
ATOM	5113		· · <del>-</del>	13.166		1.00 47.03
ATOM	5114					
			· · ·	14.370	89.364	1.00 50.25
MOTA	5115			12.779	91.565	1.00 50.94
ATOM	5116		_	13.977	91.396	1.00 51.28
MOTA	5117	CZ TYR B 26	4 59.600	14.769	90.297	1.00 52.21
ATOM	5118	OH TYR B 26	4 60.268	15.961	90.142	1.00 49.65
ATOM	5119	C TYR B 26		8.628		1.00 31.04
ATOM	5120	O TYR B 26		8.181	91.657	1.00 24.50
	5121	N LEU B 26	·			1.00 24.50
ATOM				8.074	89.666	1.00 26.68
ATOM	5122	CA LEU B 26		6.900	90.086	1.00 24.17
ATOM	5123	CB LEU B 26		6.838	89.390	1.00 24.69
MOTA	5124	CG LEU B 26		7.982	89.761	1.00 26.00
MOTA	5125	CD1 LEU B 26	5 51.970	7.583	89.460	1.00 22.21
ATOM	5126	CD2 LEU B 26	5 53.530	8.281	91.238	1.00 29.31
ATOM	5127	C LEU B 26	5 56.478	5.568	89.948	1.00 25.83
MOTA	5128	O LEU B 26		4.512	89.908	1.00 21.74
ATOM	5129	N SER B 26		5.618	89.867	
	5130	CA SER B 26		4.398		1.00 23.30
ATOM					89.813	1.00 20.75
ATOM	5131	CB SER B 26		3.900	88.378	1.00 19.67
MOTA	5132	OG SER B 26		4.615	87.739	1.00 18.11
ATOM	5133	C SER B 26		4.710	90.420	1.00 23.01
MOTA	5134	O SER B 26	60.437	5.845	90.345	1.00 17.74
ATOM	5135	N LYS B 26	7 60.590	3.707	91.023	1.00 24.25
MOTA	5136	CA LYS B 26	7 61.905	3.916	91.613	1.00 23.79
ATOM	5137	CB LYS B 26		3.153	92.929	1.00 23.71
MOTA	5138	CG LYS B 26		3.582	93.960	1.00 27.29
ATOM	5139	CD LYS B 26		5.088	94.207	
	5140					1.00 30.33
ATOM				5.535	95.273	1.00 30.90
MOTA	5141	NZ LYS B 26		7.004	95.509	1.00 33.37
ATOM	5142	C LYS B 26		3.483	90.634	1.00 26.41
ATOM	5143	O LYS B 26	64.153	3.317	91.016	1.00 25.33
MOTA	5144	N PHE B 268	62.595	3.288	89.375	1.00 22.18
ATOM	5145	CA PHE B 268	63.529	2.919	88.318	1.00 22.78
MOTA	5146	CB PHE B 268	62.814	2.171	87.179	1.00 20.55
ATOM	5147	CG PHE B 268		0.761	87.526	1.00 19.23
ATOM	5148	CD1 PHE B 268		-0.025	86.585	1.00 20.72
ATOM	5149	CD2 PHE B 268	_	0.207	88.773	1.00 18.17
	5150		-			
MOTA				-1.336	86.875	1.00 18.83
ATOM	5151	CE2 PHE B 268		-1.105	89.073	1.00 20.05
MOTA	5152	CZ PHE B 268	_	-1.879	88.122	1.0' 19.70
ATOM	5153	C PHE B 268	64.114	4.222	87.785	1.01 23.66
MOTA	5154	O PHE B 268	63.412	5.232	87.692	1.00 19.40
ATOM	5155	N ASN B 269	65.396	4.203	87.437	1.00 21.96
ATOM	5156	CA ASN B 269	66.060	5.396	86.926	1.00 25.04
ATOM	5157	CB ASN B 269	67.243	5.783	87.824	1.00 25.68
ATOM	5158	CG ASN B 269	66.845	5.946	89.273	1.00 27.04
ATOM	5159	OD1 ASN B 269	65.832		89.579	1.00 28.81
				6.557		
ATOM	5160	ND2 ASN B 269	67.659	5.419	90.176	1.00 31.12
ATOM	5161	C ASN B 269	66.579	5.151	85.523	1.00 25.87
ATOM	5162	O ASN B 269	67.769	5.336	85.268	1.00 24.58
ATOM	5163	N LEU B 270	65.695	4.757	84.611	1.00 21.37
MOTA	5164	CA LEU B 270	66.116	4.462	83.241	1.00 16.35
ATOM	5165	CB LEU B 270	65.176	3.426	82.610	1.00 24.12
ATOM	5166	CG LEU B 270	64.909	2.144	83.412	1.00 27.89
	5167	CD1 LEU B 270	64.181	1.136	82.515	1.00-23.01
ATOM		CD2 LEU B 270	66.221		83.904	1.00 23.92
ATOM	5168			1.547		
MOTA	5169	C LEU B 270	66.184	5.682	82.337	1.00 20.06
MOTA	5170	O LEU B 270	65.654	6.761	82.663	1.00 16.34
ATOM	5171	N SER B 271	66.839	5.497	81.193	1.00 20.07

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> mo>/	5172	CA	SER B	271	66.989	6.546	80.200	1.00 21.20
MOTA						6.621	79.714	1.00 21.80
MOTA	5173	CB	SER B		68.437			
MOTA	5174	OG	SER B	271	68.772	5.485	78.921	1.00 21.47
	-		SER B		66.106	6.228	79.000	1.00 22.83
ATOM	5175	С					78.854	1.00 16.12
MOTA	5176	0	SER B	271	65.631	5.102		
	5177	N	ASN B	272	65.916	7.238	78.154	1.00 20.84
MOTA					65.152	7.156	76.906	1.00 27.82
ATOM	5178	CA	ASN B				76.123	1.00 30.30
MOTA	5179	CB	ASN B	272	65.263	8.478		
	5180	CG.	ASN B	272	64.198	9.456	76.475	1.00 37.83
MOTA					64.167	10.575	75.946	1.00 37.72
ATOM	5181		ASN B					1.00 41.69
MOTA	5182	ND2	ASN B	272	63.299	9.052	77.360	
	5183	С	ASN B	272	65.701	6.088	75.974	1.00 26.88
ATOM					64.967	5.280	75.412	1.00 23.12
ATOM .	5184	0	asn b				75.774	1.00 20.40
ATOM	5185	N	VAL B	273	67.012	6.160		
	5186	CA	VAL B	273	67.745	5.260	74.899	1.00 27.34
MOTA					69.225	5.705	74.805	1.00 30.40
ATOM	5187	CB	VAL B					1.00 34.98
ATOM	5188 .	CG1	VAL B	273	70.036	4.691	74.029	
	5189	CG2	VAL B		69.299	7.057	74.115	1.00 33.57
ATOM					67.664	3.812	75.343	1.00 24.23
ATOM	5190	C	VAL B					1.00 24.19
ATOM	5191	0	VAL B	273	67.590	2.913	74.513	
	5192	N	ALA B	274	67.690	3.580	76.648	1.00 20.96
MOTA					67.589	2.220	77.151	1.00 18.12
MOTA	5193	CA	ALA B					1.00 19.09
ATOM	5194	CB	ALA .B	274	67.858	2.195	78.646	
ATOM	5195	С	ALA B	274	66.172	1.729	76.863	1.00 18.23
_			ALA B		65.962	0.567	76.525	1.00 20.77
ATOM	5196	0					77.003	1.00 18.50
MOTA	5197	N	PHE B		65.207	2.631		
ATOM	5198	CA	PHE B	275	63.802	2.310	76.761	1.00 21.25
	_		PHE B		62.941	3.546	77.037	1.00 22.24
MOTA	5199	CB				3.303	76.921	1.00 24.72
MOTA	5200	CG	PHE B		61.466			1.00 23.64
ATOM ·	5201	CD1	PHE B	275	60.815	2.483	77.826	
	5202		PHE B		60.732	3.893	75.907	1.00 27.31
MOTA					59.450	2.254	77.722	1.00 27.82
ATOM	5203		PHE B				75.795	1.00 27.62
ATOM	5204	CE2	PHE B	275	59.365	3.670		
	5205	CZ	PHE B	275	58.727	2.851	76.701	1.00 25.78
ATOM			PHE .B		63.642	1.860	75.305	1.00 24.47
MOTA	5206	C					75.030	1.00 22.68
ATOM	5207	0	PHE B	275	63.045	0.821		
ATOM	5208	N	LEU B	276	64.183	2.648	74.378	1.00 23.85
		CA	LEU B		64.128	2.330	72.946	1.00 21.28
ATOM	5209					3.421	72.134	1.00 19.87
ATOM	5210	CB	LEU B		64.814		70.662	1.00 24.94
ATOM	5211	CG	LEU B		65.114	3.132		
	5212	CD1	LEU B	276	63.818	2.852	69.936	1.00 24.81
ATOM		222	LEU B	275	65.840	4.312	70.018	1.00 21.01
MOTA	5213		LEU B	270			72.653	1.00 22.33
MOTA	5214	С	LEU B		64.841	1.021		
	5215	0	LEU B	276	64.348	0.191	71.886	1.00 20.73
ATOM			LYS B		66.011	0.857	73.261	1.00 20.72
MOTA	5216	N	ם כום	277		-0.335	73.076	1.00 24.36
ATOM	5217	CA	LYS 3		66.823			
ATOM	5218	CB	LYS B	277	68.086	-0.239	73.938	1.00 27.37
	5219	CG	LYS 3		69.303	-0.973	73.381	1.00 35.58
MOTA			LYS B		69.061	-2.456	73.188	1.00 43.87
ATOM	5220	CD					72.580	1.00 44.87
ATOM	5221	CE	LYS B		70.283	-3.137		
	5222	NZ	LYS B	277	70.616	-2.586	71.230	1.00 49.66
ATOM			LYS B	277	66.000	-1.554	73.482	1.00 24.22
MOTA	5223	С	піэ в	211		2 550	72.777	1.00 19.90
ATOM	5224	0	LYS B	277	65.987	-2.568		1.00 13.30
	5225	N	ALA B	278	65.319	-1.454	74.624	1.00 22.32
ATOM			ALA B	278	64.476	-2.544	75.114	1.00 21.71
ATOM	5226	CA					76.381	1.00 17.34
ATOM	5227	CB	ALA B		63.752	-2.117		1 00-22 60
	5228	С	ALA B	278	63.459	-2.896	74.031	1.00 22.68
ATOM		ō	ALA B		63.231	-4.068	73.723	1.00 19.27
MOTA	5229			270		-1.862	73.464	1.00 24.79
ATOM	5230	N	PHE B		62.849			1.00 22.74
	5231	CA	PHE B	279	61.860	-2.014	72.398	
ATOM		CB	PHE B		61.395	-0.629	71.955	1.00 22.46
ATOM	5232					-0.640	70.778	1.00 22.62
ATOM	5233	CG	PHE B		60.467		70.882	1.00 21.74
MOTA	5234	CD1	PHE B	279	59.196	-1.182		7.00 57.13
	5235		PHE B		60.862	-0.078	69.567	1.00 26.07
MOTA					58.325	-1.162	69.799	1.00 27.02
ATOM	5236		PHE B				68.476	1.00 25.57
MOTA	5237	CE2	PHE B	279	60.001	-0.051		1.00 20.07
ALOM.								

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MOTA	5238	CZ	PHE I	3 279	58.727	-0.594	68.592	1.00 25.13
ATOM	5239	С	PHE I	3 279	62.472	-2.768	71.212	1.00 23.60
ATOM	5240	0	PHE I			-3.697	70.678	1.00 26.54
							70.804	1.00 21.93
ATOM	5241	N	ASN I			-2.376	•	
ATOM	5242	CA	ASN I	3 280	64.318	-3.046	69.680	1.00 23.70
MOTA	5243	CB	ASN I	3 280	65.520	-2.248	69.164	1.00 22.63
							68.505	1.00 30.83
'ATOM	5244	CG	ASN I					
MOTA	5245	OD1	. ASN I	3 280			67.796	1.00 25.81
ATOM	5246	ND2	ASN B	3-280	65.900	0.112	68.714	1.00 26.54
ATOM	5247	С	ASN E	0.85	64.746	-4.466	70.009	1.00 26.10
							69.124	1.00 26.16
MOTA	5248	0	ASN I					
MOTA	5249	N	ILE E	3 281	65.080		71.272	1.00 26.10
MOTA	5250	CA	ILE E	3 281	65.485	-6.067	71.667	1.00 25.81
ATOM	5251	CB	ILE E		66.006	-6.098	73.124	1.00 28.50
					66.046		73.648	1.00 28.53
ATOM	5252	CG2						
MOTA	5253	CG1	ILE E	3 281	67.392		73 <b>.1</b> 73	
ATOM	5254	CD1	ILE E	3 281	. 68.038	-5.442	74.541	1.00 28.24
ATOM	5255	C	ILE F	281	64.320	-7.030	71.507	1.00 25.77
		ō	ILE E		64.484	-8.131	70.982	1.00 23.39
ATOM	5256							
ATOM	5257	N	VAL E		63.139	-6.618		1.00 21.30
ATOM	5258	CA	VAL E	3 282	61.961	-7.465	71.813	1.00 22.90
ATOM	5259	CB	VAL E	282	60.703	-6 <i>.</i> 775	72.387	1.00 24.07
			VAL E		59.464	-7.611	72.093	1.00 22.28
ATOM	5260							
ATOM	5261	CG2	VAL E		60.865	-6.587	73.906	1.00 26.89
MOTA	5262	С	VAL E	282	61.718	-7.795	70.339	1.00 23.87
ATOM	5263	0	VAL E	282	61.462	-8.949	69.978	1.00 22.65
ATOM	5264	N	ARG E		61.799	-6.779	69.488	1.00 23.19
						-6.971	68.060	1.00 27.95
MOTA	5265	CA	ARG E		61.576			
ATOM	5266	CB	ARG E		61.510	-5.612	67.359	1.00 25.48
ATOM	5267	CG	ARG E	283	60.337	-4.760	67.838	1.00 26.55
ATOM	5268	CD	ARG E	283	60.442	-3.333	67.339	1.00 31.52
					60.210	-3.208	65.908	1.00 24.43
MOTA	5269	NE	ARG E					
MOTA	5270	cz	ARG E		60.915	-2.414	65.116	1.00 26.45
MOTA	5271	NH1	ARG E	283	61.902	-1.676	65.622	1.00 26.04
ATOM	5272	NH2	ARG B	283	60.634	-2.356	63.825	1.00 29.64
			ARG E		62.634	-7.855	67.402	1.00 32.04
MOTA	5273	C						1.00 29.76
MOTA	5274	0	ARG B		62.341		66.431	
MOTA	5275	N	GLU B	284	63.859	-7.821	67.923	1.00 31.50
ATOM	5276	CA	GLU B	284	64.934	-8.646	67.381	1.00 32.42
	5277	CB	GLU B		66.289	-8.260	67.992	1.00 38.31
MOTA					66.798	-6.864	67.640	1.00 48.93
ATOM	5278	CG	GLU B					
ATOM	5279	CD	GLU B		68.102	-6.518	68.362	1.00 56.28
MOTA	5280	OE1	GLU B	284	69.084	-7.281	68.222	1.00 57.37
ATOM:	5281	OE2			68.150	-5.485	69.069	1.00 55.42
			GLU B		64.638	-10.105	67.714	1.00 31.93
MOTA	5282	С						1.00 28.26
ATOM	5283	0	GLU B			-11.001	66.913	
ATOM	5284	N	VAL B	285	64.089	-10.340	68.901	1.00 28.09
MOTA	5285	CA	VAL B	285	63.765	-11.697	69.325	1.00 30.67
			VAL B			-11.802	70.863	1.00 28.33
ATOM	5286	CB					71.262	1.00 29.84
ATOM	5287	CG1	VAL B	285		-13.206		
MOTA	5288	CG2	VAL B	285	65.037	-11.470	71.478	1.00 26.93
	5289	C	VAL B	285	62.460	-12.265	68.758	1.00 31.19
ATOM			1121 2	205		-13.423	68.349	1.00 31.38
ATCM	5290	0	VAL B					
ATOM	5291	N	PHE B			-11.460	68.729	1.00 28.21
MOTA	5292	CA	PHE B	.286	60.105	-11.948	68.249	1.00 25.71
	5293	CB.	PHE B		59.064	-11.853	69.374	1.00-24.57
MOTA			PHE B			-12.804	70.514	1.00 26.87
ATOM	5294	CG	PRE D	200			71.779	1.00 25.16
ATOM	5295	CD1	PHE B	286		-12.331		
ATOM	5296	CD2	PHE B	286	59.205	-14.180	70.319	1.00 22.51
	5297	CE1	PHE B	286	59.880	-13.213	72.833	1.00 22.92
ATOM.		252	PHE B	136	59.433	-15.063	71.362	1.00 21.99
ATOM	5298		rne s	200			72.626	1.00 26.75
ATOM	5299	CZ	PHE B	-50		-14.578		
ATOM	5300	С	PHE B	286		-11.318	66.993	1.00 25.90
	5301	ō	PHE B	286	58.388	-11.630	66.620	1.00 22.84
ATOM			GLY B	287		-10.451	66.329	1.00 28.27
ATOM	5302	N	GUI D	207	59.756		65.130	1.00 23.38
ATOM	5303	CA	GLY B	20/	39.130	-9.814	- 03.23	##

ATCM	5304	С	GLY B 287		58.765	-8.719	65.498	1.00 29.17
ATCM	5305		GLY B 287		58.786	-8.216	56.517	1.00 22.88
ATOM	5306		GLU B 288		57.896	-8.361	64.558	1.00 26.77
ATOM	5307		GLU B 288		56.893	-7.324	64.754	1.00 25.38
	5308		GLU B 288		56.405	-6.791	63.405	1.00 29.51
MOTA		CG			57.430			
ATCM-	5309 5310		GLU B 288			-6.003	62.605	1.00 36.06
MOTA		OE:			57.906	-4.769	63.347	1.00 41.10
ATCM	5311				57.058	-4.055	63.919	1.00 41.19
MOTA	5312	OE:			59.125	-4.503	63.348	1.00 44.69
ATOM	5313	C	-GLU B 288		55.682	-7.819	65.527	1.00 27.87
MOTA	5314	0	GLU B 288		.55.209	-8.931	65.308	1.00 26.80
MOTA	5315	N	GLY B 289		55.176	-6.973	66.419	1.00 24.53
ATOM	5316	CA	GLY B 289		54.006	-7.326	67.204	1.00 29.17
ATCM	5317	C	GLY B 289		53.015	-6.171	67.244	1.00.30.46
ATCM	5318	0	GLY B 289		53.005	-5.326	66.358	1.00 26.17
ATOM	5319	N	VAL B 290		52.171	-6.142	68.268	1.00 23.95
ATOM	5320	CA	VAL B 290		51.194	-5.079	68.440	1.00 22.25
ATOM	5321	CB	VAL B 290		49.794	-5.655	68.783	1.00 18.71
ATOM	5322		L VAL B 290		48.810	-4.525	69.047	1.00 22.67
ATOM	5323	CG2			49.289	-6.504	67.629	1.00 19.26
ATOM	5324	С	VAL B 290		51.722	-4.232	69.593	1.00 21.55
ATOM	5325	0	VAL B 290		51.960	-4.741	70.687	1.00 21.32
MOTA	5326	N	TYR B 291		51.913	-2.941	69.346	1.00 21.06
MOTA	5327	CA	TYR B 291		52.479		70.357	1.00 19.29
ATOM	5328	CB	TYR B 291		53.582	-1.216	69.711	1.00 20.40
MOTA	5329	CG	TYR B 291		54.553	-2.072	68.918	1.00 23.09
ATOM	5330		TYR B 291		54.740	-1.875	67.549	1.00 19.52
ATOM	5331	CE1			55.580	-2.712	66.809	1.00 20.67
ATOM	5332	CD2			55.234	-3.122	69.527	1.00 22.88
MOTA	5333	CE2			56.070	-3.960	68.800	1.00 26.04
ATOM ·	5334	CZ	TYR B 291		56.235	-3.752	67.442	1.00 23.44
ATOM	5335	OH	TYR B 291		57.027	-4.612	66.722	1.00 28.02
MOTA	5336	C	TYR B 291		51.465	-1.180	71.068	1.00 26.89
ATOM	5337	0	TYR B 291	•	50.668	-0.479	70.429	1.00 20.26
ATOM	5338	N	LEU B 292		51.522	-1.204	72.399	1.00 21.75
MOTA	5339	CA	LEU B 292		50.604	-0.426	73.227	1.00 22.11
ATOM	5340	CB	LEU B 292		49.765	-1.369	74.088	1.00 20.92
ATOM	5341	CG	LEU B 292		49.091	-2.542	73.375	1.00 22.94
MOTA	5342	CD1			48.328	-3.362	74.411	1.00 21.03
ATOM	5343	CD2			48.149 51.330	-2.043 0.557	72.281 74.147	1.00 18.04
ATOM	5344	C	LEU B 292 LEU B 292		52.514	0.404	74.426	1.00 21.33
ATOM	5345 5346	И О	GLY B 293		50.606	1.571	74.428	1.00 23.31
ATOM		CA	GLY B 293		51.195	2.537	75.521	1.00 20.76
ATOM	5347 5348	CA	GLY B 293		51.163	1.979	76.930	1.00 26.15
ATOM		o	GLY B 293		51.263	0.765	77.133	1.00 20.15
MOTA	5349 5350	N	GLY B 294		51.203	2.859	77.914	1.00 24.63
ATOM ATOM	5351	CA	GLY B 294		50.980	2.407	79.293	1.00 20.00
	5352	C	GLY B 294		51.176	3.538	80.285	1.00 22.59
atom atom	5353	Ö	GLY B 294		51.145	4.719	79.916	1.00 17.46
		N			51.373	3.179	81.551	1.00 17.10
ATCM	5354	CA	GLY B 295 GLY B 295		51.577	4.180	82.582	1.00 16.52
ATOM	5355		GLY B 295		52.695	5.145	82.232	1.00 10.52
ATOM	5356	C	GLY B 295				81.732	1.00 16.31
MOTA	5357 5358	O N	GLY B 295		53.738 52.467	4.737 6.430	81.732	1.00 21.93
ATOM	5358	CA N	GLY B 296		53.448	7.465	82.207	1.00 20.05
ATCM			GLY B 296		52.869	8.750	82.759	1.00 20.05
MOTA	5360	C	GLY B 296		51.790		82.739	1.00 20.48
ATOM	5361	O M	TYR B 297		53.573	9.160 9.402	83.682	1.00 20.93
ATOM	5362	И	TYR B 297		53.575	10.598	84.306	1.00 23.25
ATOM	5363	CA	TYR B 297		52.731	10.398	85.774	1.00 23.23
ATOM	5364	CB CG	TYR B 297		52.731	8.944	85.900	1.00 24.76
ATOM	5365 5366				52.779		85.936	1.00 21.97
ATOM	5366	CD1			52.779	7.758 6.514	85.912	1.00 19.79
ATOM	5367		TYR B 297		50.653		85.849	1.00 20.86
ATCM	5368	CD2				8.850	85.822	1.00 20.50
atom	5369	CE2	TYR B 297		50.012	7.612	03.042	1.00 15.57

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MOTA	5370	CZ TYR B 297	50.75	8 6.457	85.851	1.00 23.85
ATOM	5371	OH TYR B 297	50.10			
ATOM	5372	C TYR B 297	53.83			
ATOM	5373	O TYR B 297	53.45			
ATOM		N HIS B 298	54.97			,
ATOM		CA HIS B 298				
ATOM			55.78			
ATOM			57.270			
		CG HIS B 298	58.09		83.502	1.00 25.13
ATOM	5378	CD2 HIS B 298	58.406		82.482	1.00 28.22
ATOM	5379	ND1 HIS B 298	58.617	14.536	84.641	1.00 32.76
ATOM	5380	CE1 HIS B 298	59.209		84.323	
ATOM		NE2 HIS B 298	59.094	15.852	83.019	
ATOM	5382	C HIS B 298	55.589		81.795	1.00 25.66
MOTA		O HIS B 298	56.087		80.923	
MOTA	5384 1	N PRO B 299	54.901		81.496	
MOTA	5385	CD PRO B 299	54.388		82.447	1.00 29.91
MOTA	5386. (	CA PRO B 299	54.616		80.127	
ATOM	5387	CB PRO B 299	53.952		80.342	
MOTA	5388 (	G PRO B 299	54.583		81.656	
ATOM	5389	PRO B 299	55.815			1.00 27.97
ATOM	5390		55.738			1.00 27.08
MOTA	5391 N		56.925		78.057	1.00 28.58
ATOM		A TYR B 300	58.114	15.484	79.668	1.00 27.30
ATOM		B TYR B 300		15.593	78.824	1.00 27.17
ATOM		G TYR B 300	59.173	16.496		1.00 31.65
ATOM		D1 TYR B 300	58.684	17.851	79.921	1.00 31.61
ATOM			57.414	18.318	79.582	1.00 32.71
ATOM			56.971	19.568	80.014	1.00 38.52
MOTA			59.499	18.670	80.701	1.00 30.92
ATOM	5399 C		59.072	19.517	81.138	1.00 32.13
ATOM		•	57.808	20.361	80.795	1.00 39.17
ATOM			57.374	21.585	81.252	1.00 43.90
ATOM			58.731	14.218	78.572	1.00 25.20
			59.106	13.894	77.445	1.00 25.15
ATOM	5403 N 5404 C		58.845	13.419	79.628	1.00 20.55
MOTA	,		59.414	12.080	79.508	1.00 22.12
ATOM	5405 CI		59.417	11.388	80.874	1.00 17.09
ATOM	5406 C	ALA B 301	58.608	11.260	78.505	1.00 15.20
ATOM .	5407 0	ALA B 301	59.161	10.629	77.613	1.00 17.12
ATOM	5408 N		57.295	11.290	78.667	1.00 18.02
MOTA	5409 C	· · · · · · · · · · · · · · · · · · ·	56.381	10.553	77.815	1.00 19.88
MOTA	5410 CE		54.957	10.702	78.362	1.00 21.72
ATOM ·	5411 CC		53.767	10.118	77.606	1.00 31.08
ATOM		01 LEU B 302	52.576	9.980	78.549	1.00 31.35
ATOM		2 LEU B 302	53.434	11.011	76.415	1.00 27.11
ATOM	5414 C	LEU B 302	56.445	10.988	76.351	1.00 21.13
ATOM '	5415 0	LEU B 302	56.473	10.149	75.449	1.00 21.76
ATOM	5416 N		56.472	12.293	76.115	1.00 17.69
ATOM	5417 CA		56.516	12.811	74.755	1.00 17.79
ATOM	5418 CB		56.357	14.326	74.780	1.00 24.50
ATOM	5419 C	ALA B 303	57.803	12.425	74.040	1.00 20.84
MOTA	5420 O	ALA B 303	57.781	11.968	72.891	1,00 19.33
MOTA	5421 N	ARG B 304	58.930	12.594	74.723	1.00 21.08
MOTA	5422 CA	ARG B 304	60.215		74.120	1.00 25.56
MOTA	5423 CB	ARG B 304	61.375		74.962	1.00 18.37
MOTA	5424 CG	ARG B 304	61.427		75.072	1.00 23.12
MOTA	5425 CD	ARG B 304	62.797		75.624	1.00 29.00
MOTA	5426 NE	ARG B 304	63.073		76.789	1.00 29.00
MOTA	5427 CZ	ARG B 304	64.271	_		1.00 30.24
MOTA	5428 NH		65.363			
TOM		ARG B 304	64.365			1.00 24.98
MOT	5430 C	ARG B 304	60.406			1.00 36.15
TOM	5431 0	ARG B 304				1.00 20.46
TOM	5432 N	ALA B 305	60.070			1.00 18-70
TOM .	5433 CA	ALA B 305	60.226	9.988	74.937	1.00 22.48
TOM	5434 CB	ALA B 305	59.847			1.00 19.70
	5434 CB		JJ.04/			1.00 24.24
TOM	-435 C	ALA B 305	59.407	7.930	/3.711	1.00 15.82

ATOM	5436	0	ALA	B 305	59.938	7.184	72.888	1.00 19.12
MOTA	5437	N	TRP	B 306	58.113	8.230	73.659	1.00 18.65
ATOM	5438	CA	TRP	B 306	57.298	7.668	72.600	1.00 19.57
ATOM	5439	CB	TRP	B 306	55.800	7.856	72.893	1.00 18.26
ATOM	5440	CG	TRP	B 306	55.301	6.911	73.953	1.00 20.71
MOTA	5441	CD2	TRP	B 306	54.087	7.025	74.708	1.00 23.94
MOTA	5442	CE2		B 306	53.988	5.870	75.513	1.00 24.73
MOTA	5443	CE3	TRP	B 306	53.073	7.991	74.780	1.00 26.01
ATOM	5444			B 306	55.872	5.721	74.326	1.00 20.04
ATOM	5445	NE1			55.092	5.093	75.260	1.00 19.17
ATOM	5446	CZ2		в 306	52.912	5.655	76.385	1.00 28.04
MOTA	5447	CZ3	TRP	B 306	52.001	7.779	75.646	1.00 28.68
ATOM	5448	CH2		B 306	51.930	6.619	76.437	1.00 31.22
ATOM	5449	С	TRP		57.665	8.223	71.226	1.00 23.48
MOTA	5450	0	TRP	B 306	57.416	7.574	70.212	1.00 22.38
MOTA	5451	N		B 307	58.262	9.412	71.176	1.00 22.36
ATOM	5452	CA	THR	в 307	58.672	9.953	69.880	1.00 25.94
ATOM	5453	CB		B 307	59.143	11.417	69.986	1.00 25.88
ATOM	5454	OG1	THR	B 307	58.015	12.261	70.258	1.00 21.07
ATOM	5455	CG2	THR	в 307	59.827	11.864	68.686	1.00 22.52
ATOM	5456	С	THR	B 307	59.815	9.078	69.350	1.00 30.09
MOTA	5457	0	THR	B 307	59.922	8.834	68.144	1.00 25.82
ATOM	5458	N	LEU	B 308	60.664	8.596	70.258	1.00 27.54
ATOM	5459	CA	LEU	B 308	61.773	7.734	69.857	1.00 26.76
ATOM	5460	CB	LEU	B 308	62.691	7.424	71.054	1.00 24.24
MOTA	5461	CG	LEU	B 308	63.420	8.614	71.718	1.00 31.16
ATOM	5462	CD1	LEU	B 308	64.282	8.147	72.877	1.00 24.71
ATOM	5463	CD2	LEU	B 308	64.289	9.325	70.700	1.00 24.59
ATOM	5464	С	LEU	B 308	61.184	6.443	69.287	1.00 27.20
ATOM	5465	0		B 308	61.609	5.961	68.234	1.00 23.52
ATOM	5466	N	ILE	<del>-</del>	60.190	5.898	69.980	1.00 25.10
MOTA	5467	CA		B 309	59.537	4.679	69.530	1.00 25.14
ATOM	5468	CB	ILE		58.387	4.266	70.485	1.00 27.05
MOTA	5469		ILE		57.646	3.058	69.926	1.00 23.57
ATOM	5470		ILE :		58.952	3.947	71.868	1.00 22.98
ATOM	5471			В 309	59.927	2.793	71.868	1.00 24.25
ATOM	5472	C		B 309	58.958	4.885	68.133	1.00 25.41
ATOM	5473	0		B 309	59.177	4.064	67.243	1.00 22.13
MOTA	5474	N	TRP :		58.232	5.984	67.943	1.00 27.45
ATOM	5475 5476	CA		B 310	57.618	6.266	66.648	1.00 29.27
MOTA	5476 5477	CB CG		B 310 B 310	56.721	7.505	66.715	1.00 27.00
ATOM ATOM	5478			B 310	56.112 55.172	7.847	65.378	1.00 28.26
ATOM	5479			B 310	54.947	7.061 7.729	64.633	1.00 27.50
ATOM	5480			B 310	54.500	5.856	63.408 64.877	1.00 30.47 1.00 29.85
ATOM	5481			3 3 1 0 3 3 1 0	56.406	P.929	64.597	1.00 29.85 1.00 29.76
ATOM	5482			310 310	55.713	.865	63.415	1.00 25.76
MOTA	5483			3 310	54.076	234	62.429	1.00 28.23
ATOM	5484			3 310	53.636	5.362	63.901	1.00 30.24
ATOM	5485			3 310	53.433	6.053	62.692	1.00 27.63
ATOM	5486			3 310	58.629		CE 500	1.00 30.16
ATOM	5487		TRP E		58.378	5.964	64.410	1.00 30.04
ATOM	5488		CYS E		59.762	7.069	65.793	1.00 24.26
ATOM	5489		CYS E		60.782	7.233	64.764	1.00 27.97
MOTA	5490		CYS E		61.893	8.157	65.252	1.00 28.21
ATOM	5491		CYS E		61.422	9.905	65.381	1.00 33.38
ATOM	5492		CYS E		61.380	5.886	64.351	1.00 30.02
ATOM	5493		CYS E		61.670	5.660	63.172	1.00 25.45
ATOM	5494		GLU B		61.570	5.001	65.327	1.00 31.59
ATOM	5495		GLU B		62.111	3.669	65.067	1.00 33.48
ATOM	5496		GLU B		62.142	2.843	66.352	1.00 34.78
MOTA	5497		GLU B		63.487	2.307	66.758	1.00 39.45
MOTA	5498		GLU B	312	64.171	1.513	65.675	1.00 40.11
ATOM .	5499	0E1 :	GLU B		63.539	0.614	65.081	1.00 43.69
ATOM	5500		GLU B		65.358	1.782	65.437	1.00 39.26
ATCM	5501	C	GLU B	312	61.197	2.959	64.080	1.00 29.97

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#### Figure 18-84.

ATOM	5502	. 0	.GLU B	312		61.640	2.49	97 63.03	RS 10	0 31.38
MOTA	5503		LEU B			59.919				0 26.70
MOTA	5504	CA	LEU B	313		58.930	2.20			0 26.73
MOTA	5505			313		57.571	2.17	73 64.29		0 25.83
MOTA	5506			313		57.429				0 35.18
MOTA	5507			313		56.063				0 32.49
ATOM	5508			313		57.595				0 29.71
MOTA	5509	C		313		58.768		_ '		0 29.03
MOTA	5510			313		58.716				0 25.39
MOTA MOTA	5511 5512	N CA		314		58.677 58.498				0 30.13
ATOM	5513	CB	SER B	314	-	58.206				0 34.06
ATOM	5514	OG	SER B			57.041				0 31.15 0 48.58
ATOM	5515	c	SER B			59.707				0 31.84
ATOM	5516	0	SER B			59.632		-		0 34.15
ATOM	5517	N	GLY B			60.831				31.81
MOTA	5518	CA	GLY B	315		62.036	4.48			37.27
MOTA	5519	С	GLY B	315		62.659	5.85			39.93
MOTA	5520	0	GLY B			63.363	6.05		4 1.00	39.79
ATOM	5521	N	ARG B			62.422				38.22
MOTA	5522	CA	ARG B			63.004	8.12			38.66
ATOM	5523 5524	CB CG		316		61.908	9.18			40.20
MOTA MOTA	5525	CD	ARG B ARG B			61.089 60.032	9.34 10.39			39.00
ATOM	5526	NE	ARG B			59.002	9.95			42.13 45.09
ATOM	5527	CZ	ARG B			58.075	10.75			40.84
ATOM	5528	NH1				58.064	12.03			48.44
ATOM	5529	NH2	ARG B	316		57.150	10.27			35.96
MOTA	5530	C	ARG B			64.031	8.46			39.03
ATOM	5531	0	ARG B			63.952	7.98			34.34
ATOM	5532	N	GLU B			65.003	9.29			39.58
MOTA MOTA	5533 5534	CA CB	GLU B			66.074 67.142	9.69			43.35
ATOM	5535	CG	GLU B			67.609	10.50 9.91			) 49.34 ) 57.04
ATOM	5536	CD	GLU B			66.546	10.00			62.79
ATOM	5537		GLU B			66.146	11.14			63.46
ATOM	5538	OE2	GLU B	317		66.108	8.95	4 58.28		64.46
MOTA	5539	С		317		65.555	10.52			41.58
ATOM	5540	0		317		64.658	11.35			39.74
ATOM	. 5541	N		318		66.118	10.30			35.38
MOTA MOTA	5542 5543	CA CB	VAL B 3			65.706 66.000	11.04			38.76
ATOM	5544		VAL B 3			65.560	10.26			42.28 38.26
ATOM	5545					65.287	8.91			39.99
ATOM	5546	C	VAL B 3			66.459	12.37			41.82
.TOM	5547	0	VAL B 3	318		67.689	12.39			37.20
MOTA	5548	N	PRO B 3			65.735	13.49	1 65.356	1.00	43.18
MOTa	5549	CD	PRO B 3			64.290	13.672			41.90
ATOM	5550	CA	PRO B 3			66.402	14.792			44.31
MOTA	5551 5552	CB CG	PRO B 3			65.241 64.079	15.763			44.58
MOTA MOTA	5553	C	PRO B 3			67.086	15.011 14.96			43.34
ATOM	5554	ŏ	PRO B 3			66.541	14.565			44.62 43.75
ATOM	.5555	N	GLU B 3			68.277	15.552			44.16
ATOM	5556	CA	GLU B 3			69.029	15.762			45.92
MOTA	5557	CB	GLU B 3	20	•	70.381	16.406			50.87
ATOM	5558	CG	GLU B 3			71.165	16.768		1.00	53.71
MOTA	5559	CD	GLU B 3			72.455	17.505			57.75
MOTA	5560		GLU B 3			73.161	17.874			56.37
ATOM	5561	OE2	GLU B 3.			72.762	17.714			60.07
ATOM	5562	C	GLU B 3			68.311	16.625			44,42
ATOM	5563 5564	O M	GLU B 3			68.244 67.778	16.279			42.32
MOTA MOTA	5565	N CA	LYS B 3:			57.102	17.753 18.672			42.50 45.24
MOTA	5566	CB	LYS B 3			57.853	20.000			46.43
MOTA	5567	CG	LYS B 3			57.890	20.802			51.45
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ATOM	5568	CD	LYS	B 321	. (	58.700	20.144	67.057	1.00 57.24
ATOM	5569	CE	LYS	B 321	. (	57.936	19.062	66.280	1.00 55.24
MOTA	5570	NZ	LYS	B 321	. (	66.738	19.588	65,558	1.00 55.31
MOTA	5571	С		B 321		55.662	18.971	69.098	1.00 43.44
MOTA	5572	0	LYS	B 321	. (	55.211	18.736	67.978	1.00 43.03
ATOM	5573	N	LEU	B 322	. 6	54.947	19.512	70.076	1.00 39.45
ATOM	5574	CA	LEU	B 322	: 6	53.563	19.885	69.875	1.00 40.31
ATOM	5575		LEU			2.846	20.034	71.215	1.00 40.88
ATOM	5576		LEU		: 6	52.943	18.901	72.234	1.00 40.09
ATOM	5577			B 322		2.001	19.175	73.388	1.00 38.17
ATOM	5578		LEU			2.588	17.596	71.580	1.00 41:56
ATOM	5579	С	LEU			3.615	21.244	69.197	1.00 41.23
ATOM	5580	0		B 322		4.466	22.070	69.531	1.00 39.22
ATOM	5581	N		B 323		2.735	21.473	68.233	1.00 40.04
ATOM	5582	CA		B 323		2.703	22.771	67.582	1.00 43.32
. ATOM	5583	CB		B 323		1.985	22.707	66.234	1.00 41.53
ATOM	5584	CG		B 323		0.617	22.085	66.335	1.00 41.89
ATOM	5585	OD1	ASN	B 323	5	9.889	22.308	67.304	1.00 39.79
ATOM	5586	ND2	ASN	B 323	6	0.243	21.317	65.317	1.00 40.43
'ATOM	5587	С	ASN	B 323	ε	1.949	23.690	68.532	1.00 44.76
ATOM	5588	0	ASN	B 323	6	1.402	23.237	69.539	1.00 45.80
MOTA	5589	N	ASN	B 324	$\epsilon$	1.902	24.973	68.210	1.00 46.85
ATOM	5590	CA		B 324		1.234	25.930	69.076	1.00 47.60
ATOM	5591	CB	ASN	B 324	6	1.460	27.348	68.549	1.00 50.87
ATOM	5592	CG		B 324	6	1.089	28.407	69.562	1.00 55.06
ATOM	5593			B 324	5	9.925	28.565	69.919	1.00 60.68
MOTA	5594	ND2	ASN	B 324	6	2.091	29.131	70.048	1.00 59.17
ATOM	5595	С	ASN	B 324	5	9.740	25.664	69.249	1.00 43.97
ATOM	5596	0	ASN	B 324		9.190	25.898	70.322	1.00 41.33
MOTA	5597	N	LYS			9.087	25.168	68.201	1.00 43.49
ATOM	5598	CA	LYS			7.655	24.892	68.264	1.00 45.95
ATOM	5599	CB		B 325		7.112	24.415	66.909	1.00 48.97
ATOM	5600	CG	LYS			7.212	25.400	65.731	1.00 53.41
ATOM	5601	CD	LYS			8.582	25.386	65.024	1.00 58.77
MOTA	5602	CE	LYS			9.700	26.013	65.846	1.00 58.10
MOTA	5603	NZ	LYS			1.024	25.906	65.178	1.00 53.38
ATOM	5604	C	LYS			7.368	23.822	69.309	1.00 45.79
MOTA	5605	0	LYS			6.375	23.891	70.034 69.381	1.00 43.91
ATOM	5606	N	ALA			8.245	22.829	70.336	1.00 44.28 1.00 44.25
ATOM	5607	CA	ALA			8.078	21.746 20.589		1.00 41.44
ATOM	5608	CB	ALA ALA			9.013 8.342	22.233	71.757	1.00 40.92
ATOM	5609	C	ALA			7.639	21.843	72.688	1.00 39.02
ATOM	5610	NO	LYS			9.352	23.085	71.922	1.00 39.02
ATOM	5611 5612	CA	LYS			9.689	23.603	73.246	1.00 40.11
ATOM	5613	CB	LYS			0.892	24.552	73.178	1.00 42.36
ATOM	5614	CG	LYS			2.174	23.922	72.(59	1.00 45.78
atom atom	5615	CD	LYS			3.325	24.926	72.675	1.00 48.46
ATOM	5616	CE		B 327		4.594	24.367	72.031	1.00 49.62
ATOM	5617		LYS			5:108	23.139	72.700	
ATOM	5618	c		B 327		8.500	24.338	73.841	1.00 39.17
ATOM	5619	ŏ		B 327		8.132	24.112	74.994	1.00 38.87
ATOM	5620	N		B 328		7.898	25.215	73.048	1.00 41.06
ATOM	5621	CA		B 328		6.750	25.986	73.512	1.00 42.35
ATOM	5622	CB	GLU I	B 328		6.357	27.028	72.463	1.00 44.02
ATOM	5623	CG		328		7.434	28.084	72.258	1.00 44.80
ATOM	5624	CD	GLU I	328		7.835	28.742	73.569	1.00 48.40
ATOM	5625		GLU I	328		6.949	29.317	74.237	1.00 51.20
ATOM	5626		GLU 1			9.029	28.680	73.935	1.00 47.81
ATOM	5627	С		3 328		5.569	25.087	73.839	1.00 38.67
ATOM	5628	9		3 3 2 8		4.794	25.377	74.750	1.00 41.20
ATOM	5629	N	LEU I			5.429	23.999	73.090	1.00 35.31
ATOM	5630	CA	LEU !			4.349	23.056	73.334	1.00 32.69
ATOM.	5631	CB		3 329		4.404	21.900	72.334	1.00 35.06
ATOM	5632	CG	LEU I	3 329	5	3.344	20.813	72.544	1.00 35.01
TOM	5633	CD1	LEU I	3 3 2 9		1.958	21.430	72.419	1.00 36.90
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MOTA	5634	CD	2 LEU	B 329		53.521	19.699	71.525	1.00 32.36
MOTA	5635		LEU			54.504		74.747	1.00 34.07
ATOM	5636			B 329		53.621	22.664	75.583	1.00 30.53
ATOM	5637			B 330		55.640	21.873	75.013	1.00 30.33
ATOM	5638			B 330		55.889		76.330	
				B 330			21.311		1.00 34.99
MOTA	5639					57.267	20.642	76.382	1.00 37.01
ATOM	5640			В 330		57.466	19.428	75.470	1.00 34.91
ATOM	5641			B 330		58.832	18.817	75.728	1.00 34.69
ATOM	5642	CD				56.369	18.396	75.742	1.00 34.10
ATOM	5643	С		B 330		55.789	22.363	77.429	1.00 37.12
ATOM	5644	Ó	LEU	В 330		55.210	22.110	78.482	1.00 34.19
ATOM	5645	N	LYS	B 331		56.353	23.540	77.186	1.00 34.34
ATOM	5646	CA	LYS	B 331		56.313	24.604	78.181	1.00 43.35
ATOM	5647	CB	LYS	B 331		57.162	25.788	77.712	1.00 46.25
ATOM	5648	CG		B 331		58.658	25.496	77.685	1.00 51.07
ATOM	5649	CD		B 331		59.482	26.610	77.021	1.00 49.96
ATOM	5650	CE		B 331		59.371	27.957	77.733	1.00 53.08
ATOM	5651	NZ		B 331		58.013	28.569	77.662	1.00 56.18
MOTA	5652	C		B 331		54.892	25.069	78.494	1.00 42.06
ATOM	5653	Ō.		B 331		54.588	25.416	79.631	1.00 43.05
	5654	N.		B 332					
ATOM						54.018	25.056	77.492	1.00 44.54
MOTA	5655	CA		B 332		52.639	25.502	77.679	1.00 46.58
ATOM	5656	CB		B 332		51.975	25.751	76.329	1.00 48.75
ATOM	5657	OG		B 332		51.769	24.527	75.646	1.00 49.55
ATOM	5658	C		B 332		51.780	24.507	78.451	1.00 49.56
ATOM	5659	0		B 332		50.618	24.791	78.749	
ATOM	5660	N			• =	52.341	23.345	78.770	1.00 50.55
MOTA	5661	CA	ILE			51.586	22.326	79.488	1.00 51.93
ATOM	5662	CB	ILE			52.259	20.945	79.376	1.00 51.82
MOTA	5663	CG2				51.447	19.902	80.134	1.00 50.29
MOTA	5664	CG1				52.359	20.539	77.905	1.00 52.18
ATOM	5665	CD1		B 333		53.044	19.210	77.693	1.00 55.42
ATOM	5666	C	ILE	B 333	•	51.367	22.634	80.964	1.00 51.45
MOTA	5667	0	ΙĻΕ	B 333		52.180	23.290	81.614	1.00 50.96
ATOM	5668	N	ASP	B 334		50.245	22.141	81.472	1:00 54.05
ATOM	5669	CA	ASP	B 334		49.850	22.306	82.865	1.00 58.15
ATOM	5670	CB	ASP	B 334		48.320	22.216	82.959	1.00 60.38
ATOM	. 5671	CG	ASP	B 334		47.751	20.972	82.262	1.00 63.85
ATOM	5672	OD1	ASP	B 334		48.017	19.833	82.710	1.00 59.16
ATOM	5673			B 334		47.033	21.138	81.252	1.00 59.71
ATOM	5674	С		B 334		50.506	21.207	83.701	1.00 55.47
MOTA	5675	0		B 334		49.833	20.291	84.171	1.00 54.08
ATOM	5676	N		B 335		51.816	21.307	83.906	1.00 54.60
MOTA	5677	CA		B 335		52.524	20.266	84.641	1.00 56.60
ATOM	5678	СВ		B 335		53.718	19.784	83.811	1.00 53.01
ATOM	5679	CG		B 335		54.522	18.717	84.482	1.00 49.30
MOTA	5686			B 335		53.898	17.589	85.008	1.00 45.61
ATOM	5681		PHE			55.901	18.843	84.605	1.00 46.83
ATOM	5682		PHE			54.637	16.600	85.651	1.00 45.95
ATOM	5683			B 335				85.247	1.00 46.02
	5684					56.651	17.860		
MOTA		CZ		B 335		56.018	16.737	85.772	1.00 46.08
ATOM	5685	C		B 335		52.971	20.559	86.072	1.00 57.29
ATOM	5686	0		B 335		52.197	20.378	87.012	1.00 63.54
MOTA	5687	N		B 336		54.223	20.983	86.229	1.00 55.21
MOTA	5688			B 336		54.818	21.286	87.535	1.00 60.30
MOTA	5689			B 336		53.783	21.846	88.517	1.00 64.95
ATOM	5690			B 336		54.375	22.225	89.867	1.00 -71.50
ATOM	5691			B 336		53.363	22.882	90.787	1.00 75.37
ATOM	5692	OE1	GLU 1	B 336		52.796	23.925	90.394	1.00 75.32
ATOM	5693	OE2	GLU I	336		53.137	22.361	91.901	1.00 76.84
ATOM	5694		GLU I			55.485	20.058	88.146	1.00 55.66
ATOM	5695		GLU E	3 3 3 6		54.823	19.093	88.529	1.00-49.97
ATOM	5696		GLU E			56.807	20.125	88.240	1.00 54.26
ATOM ·	5697		GLU I			57.630	19.047	88.767	1.00 54.35
ATOM	5698		GLU E			59.101	19.457	88,635	1.00 54.08
ATOM	5699		GLU E			60.074	18.315	88.514	1.00 54.15
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ATOM	5700	CD GLU B 337	.59.856	17.496	87.259	1.00 48.94
MOTA	5701		59.958		86.142	1.00 41.06
MOTA	5702	OE2 GLU B 337	59.581		87.391	1.00 50.23
ATOM	5703		57.278	18.740	90.227	1.00 55.08
ATOM	5704	O GLU B 337	57.130		91.039	1.00 54.51
ATOM	5705	N PHE B 338	57.140	17.458	90.557	1.00 56.20
ATOM	5706		56.798	17.048	91.918	1.00 57.73
MOTA	5707		56.713	15.527	92.020	1.00 58.47
ATOM	5708		56.231	15.034	93.359	1.00 63.17
MOTA	5709		54.882	15.096	93.696	1.00 64.86
ATOM	5710		57.129	14.526	94.293	1.00 63.51
MOTA	5711	CE1 PHE B 338	54.434	14.656	94.943	1.00 65.25
ATOM	5712	CE2 PHE B 338	56.693	14.087	95.539	1.00 63.64
MOTA	5713	CZ PHE B 338	55.342	14.152	95.864	1.00 66.30
MOTA	5714	C PHE B 338	57.836	17.539	92.918	1.00 61.85
MOTA	5 <b>71</b> 5	O PHE B 338	57.520	17.807	94.078	1.00 58.15
MOTA	5716	N ASP B 339	59.081	17.636	92.466	1.00 64.63
MOTA	5717	CA ASP B 339	60.167	18.099	93.316	1.00 67.53
MOTA	5718	CB ASP B 339	61.286	17.059	93.362	1.00 67.32
ATOM ·	5719	CG ASP B 339	62.474	17.524		1.00 68.13
MOTA	5720	OD1 ASP B 339	62.280	17.909	95.346	1.00 68.68
MOTA	5721	OD2 ASP B 339	63.603	17.502	93.646	1.00 69.03
ATOM	5722	C ASP B 339	60.718	19.435	92.829	1.00 69.03
ATOM	5723	O ASP B 339	61.211	19.545	91.708	1.00 67.54
ATOM	5724	N ASP B 340	60.626	20.442	93.693	1.00 72.19
MOTA	5725	CA ASP B 340	61.088	21.797	93.402	1.00 75.20
ATOM	5726	CB ASP B 340	61.113	22.623	94.689	1.00 77.04
ATOM	5727	CG ASP B 340	59.766	22.671	95.375	1.00 78.70
ATOM	5728	OD1 ASP B 340	58.803	23.181	94.763	1.00 79.66
ATOM	5729	OD2 ASP B 340	59.668	22.194	96.525	1.00 80.53
ATOM	5730	C ASP B 340	62.464	21.856	92.751	1.00 74.82
ATOM	5731	O ASP B 340	62.615	22.400	91.659	1.00 78.48
ATOM	5732	N GLU B 341	63.465	21.303	93.426	1.00 74.11
MOTA	5733 5734	CA GLU B 341 CB GLU B 341	64.827	21.312	92.907	1.00 76.25
MOTA	5735		65.818	21.596	94.040	1.00 79.54
ATOM ATOM	5736	CG GLU B 341 CD GLU B 341	67.277 67.539	21.653	93.596	1.00 82.33
ATOM	5737	OE1 GLU B 341	67.333	22.750 23.937	92.577 92.910	1.00 83.24 1.00 85.25
ATOM	5738	OE2 GLU B 341	67.950	22.427	91.443	1.00 83.23
ATOM	5739	C GLU B 341	65.196	19.998	92.227	1.00 83.72
ATOM	5740	O GLU B 341	65.627	19.051	92.883	1.00 77.10
ATOM	5741	N VAL B 342	65.033	19.946	90.910	1.00 77.10
ATOM	5742	CA VAL B 342	65.354	18.744	90.151	1.00 68.51
ATOM	5743	CB VAL B 342	64.081	18.027	89.663	1.00 68.83
ATOM	5744	CG1 VAL B 342	63.268	17.552	90.837	1.00 67.57
ATOM	5745	CG2 VAL B 342	63.255	18.969	88.806	1.00 67.72
ATOM	5746	C VAL B 342	66.201	19.059	88.927	1.00 65 35
ATOM	5747	O VAL B 342	67.177	18.366	88.640	1.00 68.31
MOTA	5748	N ASP'B 343	65.819	20.112	88.213	1.00 60.89
MOTA	5749	CA ASP B 343	66.514	20.520	86.998	1.00 58.89
MOTA	5750	CB ASP B 343	68.024	20.636	87.223	1.00 63.48
ATOM	5751	CG ASP B 343	68.763	21.070	85.966	.1.00 66.69
ATOM	5752	OD1 ASP B 343	70.012	21.070	85.970	1.00 67.64
ATOM	5753	OD2 ASP B 343	68.089	21.420	84.973	1.00 65.42
ATOM	5754	C ASP B 343	66.264	19.499	85.900	1.00 53.17
ATOM	5755	O ASP B 343	66.993	18.516	85.766	1.00 49.70
MOTA	5756	N ARG B 344	65.216	19.735	85.124	1.00 50.24
MOTA	5757	CA ARG B 344	64.868	18.853	84.022	1.00 46.49
ATOM	5758	CB ARG B 344	63.467	18.269	84.228	1.00 42.41
MOTA	5759	CG ARG B 344	63.317	17.367	85.452	1.00 38.59
ATOM	5760	CD ARG B 344	64.344	16.246	85.432	1.00 37.12
ATOM	5761 =762	NE ARG B 344 CZ ARG B 344	64.169	15.310	86.537	1.00 36.55
MOTA	5762	CZ ARG B 344 NH1 ARG B 344	65.078	14.413	86.905	1.00 37.20
ATOM .	5763 5764	NH2 ARG B 344	66.234 64.830	14.331	86.259 87 015	1.00 33.53
ATOM	5764 =765	C ARG B 344		13.595	87.915 82.732	1.00 28.79
MOTA	5765	~ W/G D 244	64.910	19.660		1.00 44.45

ATOM	576€	6 O ARG B 344	64.328	19.269	81.720	1.00 38.73
ATOM	5767	N SER B 345	65.618			
ATOM	5768	CA SER B 345	65.740			
MOTA	5769	CB SER B 345	66.661			
ATOM	5770	OG SER B 345	67.956			
ATOM			66.244			
ATOM	5772	= =				
ATOM	5773		65.840	21.333		
ATOM	5774		67.117			
			67.661	19.264		
MOTA	5775		68.660			
ATOM	5776		68.054	17.146	80.774	
MOTA	5777		67.433	16.013	80.240	1.00 37.62
ATOM	5778		66.843	15.048	81.077	1.00 36.73
ATOM	5779		68.072	17.294	82.157	1.00 36.40
MOTA	5780		67.489	16.344	82.999	1.00 36.54
ATOM	5781	CZ TYR B 346	66.878	15.228	82.457	1.00 36.54
ATOM	5782	OH TYR B 346	66.310	14.306	83.306	1.00 33.35
ATOM	5783	C TYR B 346	66.563	18.599	78.570	1.00 36.26
ATOM	5784	O TYR B 346	66.719	18.385	77.367	1.00 40.50
ATOM	5785	N MET B 347	. 65.445	18.282	79.214	1.00 40.30
ATOM	5786	CA MET B 347	64.346	17.628	78.516	1.00 35.43
ATOM	5787	CB MET B 347	63.280		79.513	
ATOM	5788	CG MET B 347	63.819	16.292	80.635	1.00 34.36
ATOM	5789	SD MET B 347	62,515	15.604	81.669	1.00 28.32
ATOM	5790	CE MET B 347	61.654			1.00 34.47
ATOM	5791	C MET B 347	63.701	17.027	82.142	1.00 39.60
ATOM	5792	O MET B 347		18.525	77.465	1.00 39.04
ATOM	5793	N LEU B 348	63.060	18.029	76.540	1.00 37.38
ATOM	5794	CA LEU B 348	63.857	19.839	77.606	1.00 39.21
ATOM	5795	CB LEU B 348	63.272	20.773	76.645	1.00 40.81
ATOM	5796	CG LEU. B 348	62.806	22.058	77.339	1.00 36.87
MOTA	5797	CD1 LEU B 348	61.690	21.975	78.384	1.00 42.66
ATOM	5798		61.507	23.337	79.032	1.00 43.41
ATOM	5799		60.391	21.511	77.741	1.00 40.47
ATOM	5800		64.289	21,133	75.573	1.00 41.13
ATOM	5801		64.018	21.968	74.711	1.00 38.93
	5802	N GLU B 349	65.455	20.495	75.632	1.00 37.70
ATOM		CA GLU B 349	66.527	20.757	74.681	1.00 42.48
ATOM.	5803 5804	CB GLU B 349	67.856	20.953	75.422	1.00 45.02
ATOM		CG GLU B 349	67.834	22.035	76.493	1.00 53.82
ATOM	5805	CD GLU B 349	67.483	23.402	75.938	1.00 57.46
ATOM	5806	OE1 GLU B 349	68.211	23.885	75.044	1.00 59.62
MOTA	5807	OE2 GLU B 349	66.480	23.993	76.397	1.00 57.91
MOTA	5808	C GLU B 349	66.709	19.638	73.664	1.00 43.57
ATOM	5809	O GLU B 349	66.577	19.849	72.459	1.00 41.26
ATOM	5810	N THR B 350	67.02 <b>7</b>	18.448	74.161	1.00 41.95
ATOM		CA THR B 350	67.264	17.299	73.298	1.00 40.02
ATOM	5812	CB THR B 350	68.689	16.775	73.504	1.00 43.08
MOTA	5813	OG1 THR B 350	68.894	16.490	74.894	1.00 41.07
ATOM .	5814	CG2 THR B 350	69.703	17.816	73.049	1.00 45.05
ATOM	5815	C THR B 350	66.278	16.154	73.510	1.00 37.56
ATOM	5816	O THR B 350	65.754	15.966	74.611	1.00 33.64
ATOM	5817	N LEU B 351	66.043	15.391	72.445	1.00 32.86
MOTA	5818	CA LEU B 351	65.126	14.260	72.475	1.00 35.00
ATOM	5819	CB LEU B 351	64.776	13.810	71.053	1.00 31.61
ATOM	5820	CG LEU B 351	63.709	14.601	70.312	1.00 35.31
MOTA	5821	CD1 LEU B 351	63.552	14.064	68.904	1.00 37.88
ATOM	5822	CD2 LEU B 351	62.397	14.474	71.068	1.00.39.36
MOTA	5823	C LEU B 351	65.662	13.065	73.240	1.00 33.33
ATOM	5824	O LEU B 351	64.956	12.469	74.046	1.00 31.48
ATOM	5825	N LYS B 352	66.915	12.720	72.981	1.00 29.58
ATOM		CA LYS B 352	67.527	11.576	73.633	1.00 36.77
ATOM		CB LYS B 352		10.864	72.647.	1.00 34.32
ATOM .		CG LYS B 352		10.563	71.326	1.00 39.29
MOTA		CD LYS B 352	68.703	9.949	70.294	1.00 42.25
ATOM		CE LYS B 352	69.110	8.541	70.655	1.00 46.22
ATOM		NZ LYS B 352	69.831	7.905	69.516	1.00 44.15
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### Figure 18-89

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MOTA	5832	C LYS B 352	68.295	11.983	74.878	1.00 36.30
ATOM	5833		69.086			
ATOM	5834		68.049			
MOTA	5835		68.757		77.188	1.00 33.99
ATOM	5836		67.852		78.394	1.00 38.57
ATOM	5837		67.134		78.315	1.00 43.90
ATOM	5838		66.034		78.926	1.00 22.39
ATOM	5839		67.679		77.649	1.00 50.42
ATOM	5840	C ASP B 353	70.022		77.202	
ATOM	5841	O .ASP B 353	70.189		76.368	1.00 35.83
ATOM	5842	N PRO B 354	70.954		78.116	1.00 23.71
ATOM	5843	CD PRO B 354	70.928		79.132	1.00 38.38
ATOM	5844	CA PRO B 354	72.205		78.212	
ATOM	5845	CB PRO B 354	73.003		79.213	1.00 33.62 1.00 34.46
ATOM	5846	CG PRO B 354	71.896	11.556	80.164	1.00 34.48
ATOM	5847		71.924	8.883	78.733	1.00 38.08
ATOM	5848	O PRO B 354	70.894		79.366	
ATOM	5849	N TRP B 355	72.833	7.954	78.468	1.00 24.82 1.00 31.76
ATOM	5850	CA TRP B 355	72.635	6.611	78.969	
ATOM	5851	CB TRP B 355	73.653	5.655	78.359	1.00 30.01
ATOM	5852	CG TRP B 355	73.025	4.378	77.910	1.00 34.02 1.00 44.37
MOTA	5853	CD2 TRP B 355	73.263	3.072	78.436	
ATOM	5854	CE2 TRP B 355	72.418	2.177	77.734	1.00 45.39
ATOM	5855	CE3 TRP B 355	74.107	2.569	79.432	1.00 44.31
ATOM	5856	CD1 TRP B 355	72.073	4.230	76.935	1.00 47.19 1.00 42.18
ATOM	5857	NE1 TRP B 355	71.704	2.910	76.826	1.00 37.84
ATOM	5858	CZ2 TRP B 355	72.395	0.808	77.999	1.00 37.84
ATOM	5859	CZ3 TRP B 355	. 74.084	1.207	79.694	1.00 50.83
ATOM	5860	CH2 TRP B 355	73.231	0.341	78.979	1.00 30.83
MOTA	5861	C TRP B 355	72.819	6.685	80.485	1.00 30.87
ATOM	5862	O TRP B 355	73.622	7.474	80.981	1.00 26.93
ATOM	5863	N ARG B 356	72.061	5.880	81.218	1.00 24.96
ATOM	5864	CA ARG B 356	72.147	5.848	82.671	1.00 23.57
ATOM	5865	CB ARG B 356	70.811	6:319	83.257	1.00 24.71
ATOM	5866	CG ARG B 356	70.534	7.795	82.941	1.00 23.66
ATOM	5867	CD ARG B 356	69.067	8.212	83.055	1.00 20.14
ATOM	5868	NE ARG B 356	68.926	9.610	82.642	1.00 20.59
ATOM	5869	CZ ARG B 356	67.787	10.192	82.288	1.00 25.41
ATOM	5870	NH1 ARG B 356	66.644	9.508	82.287	1.00 17.01
ATOM	5871	NH2 ARG B 356	67.796	11.464	81.910	1.00 20.07
ATOM	5872	C ARG B 356	72.481	4.410	83.085	1.00 26.57
ATOM	5873	O ARG B 356	71.610	3.641	83.485	1.00 23.02
ATOM	5874	N GLY B 357	73.761	4.063	82.978	1.00 23.92
ATOM	5875	CA GLY B 357	74.186	2.712	83.294	1.00 25.54
ATOM	5876	C GLY B 357	74.796	2.464	84.657	1.00 24.35
ATOM	5877	0 · GLY B 357	74.523.	3.161	85.628	1.00 25.88
ATCM	5878	N GLY B 358	75.638	1.444	84.718	1.00 24.32
ATOM	5879	CA GLY B 358	76.282	1.070	85.960	1.00 23.56
ATCM	5880	C GLY B 358	76.412	-0.441	85.924	1.00 29.26
ATOM	5881	O GLY B 358	76.146	~1.051	84.889	1.00 23.71
ATOM	5882	N GLU B 359	76.814	-1.051	87.033	1.00 27.64
ATOM	5883	CA GLU B 359	76.955	~2.503	87.078	1.00 32.16
ATOM	5884	CB GLU B 359	77.822	-2.936	88.265	1.00 30.40
ATOM	5885	CG GLU B 359	77.125	-2.772	89.601	1.00 31.23
ATOM	5886	CD GLU B 359	77.844	-3.479	90.741	1.00 37.96
ATCM	5887	OE1 GLU B 359	77.287	~3.521	91.861	1.00 33.89
ATOM	5888	OE2 GLU B 359	78.959	~3.990	90.520	1.00 37.40
ATOM	5889	C GLU B 359	75.571	-3.122	87.261	1.00 31.35
ATOM		O GLU B 359	74.612	-2.429	87.588	1.00 25.15
ATOM		N VAL B 360	75.482	-4.428	87.053	1.00 29.61
ATCM		CA VAL B 360	74.230	-5.147	87.251	1.00 26.21
ATOM		CB VAL B 360	74.035	-6.270	86.200	1.00 28.47
ATOM		CG1 VAL B 360	72.764	-7.045	86.492	1.00 22.74
ATOM	_	CG2 VAL B 360	73.969	-5.670	84.796	1.00 29.70
ATOM	_	C VAL B 360	74.342	-5.784	88.625	1.00 26.00
ATCM	5897	O VAL B 360	75.150	-6.693	88.821	1.00 27.55
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# **206/263** Figure 18-90

MOTA	5898	N ARG B 3	61 73.553	-5.289	89.575	1.00 26.45
ATOM	5899				90.935	
						1.00 28.47
ATOM	5900	CB ARG B 3			91.787	1.00 30.55
ATOM	5901	CG ARG B 3	61 72.937	7 -3.877	92.485	1.00 32.61
ATOM	5902	CD ARG B 3			93.117	1.00 35.00
ATOM	5903	NE ARG B 3	61 70.858	~2.617	92.094	1.00 30.31
ATOM	5904	CZ ARG B 3	61 69.753	-1.925	92.350	1.00 29.45
ATOM	5905			-1.689		
ATOM	5906				91.348	1.00 30.49
MOTA	5907	C ARG B 3	61 73.351	7.322	91.001	1.00 30.17
ATOM	5908	O ARG B 3			90.168	1.00 23.60
MOTA	5909	N LYS B 3			92.022	
MOTA	5910	CA LYS B 3	62 73.864	-9.351	92.272	1.00 36.94
ATOM	5911	CB LYS B 3	62 74.687	-9.706	93.513	1.00 40.24
ATOM	5912	CG LYS B 3		– –	93.337	1.00 52.55
ATOM	5913	CD LYS B 3	62 76.571	-8.126	92.849	1.00 56.65
MOTA .	5914	CE LYS B 3	62 76.149	-7.032	93.819	1.00 53.39
		NZ LYS B 3				
ATOM	5915				93.341	1.00 48.87
ATOM	5916	C LYS B 3	<b>52</b> 72.427	-9.826	92.463	1.00 32.84
MOTA	5917	O LYS B 3	52 72.045	-10.867	91.938	1.00 28.27
MOTA	5918				93.215	1.00 34.67
ATOM	5919	CA GLU B 3			93.435	1.00 35.72
ATOM	5920	CB GLU B 3	69.519	-8.532	94.390	1.00 36.04
	5921	CG GLU B 3		-7.077	93.977	1.00 44.81
ATOM						
ATOM	5922	CD GLU B 3		-6.186	95.033	1.00 52.14
ATOM	5923	OE1 GLU B 3	67.661	-6.370	95.341	1.00 48.46
ATOM	5924	OE2 GLU B 3		-5.300	95.566	1.00 57.31
MOTA	5925	C GLU B 3		-9.619	92.111	1.00 30.68
ATOM	5926	O GLU B 3	53 68.695	-10.530	91.944	1.00 30.45
ATOM	5927	N VAL B 3	69.784	-8.724	91.166	1.00 26.19
ATOM	5928	CA VAL B 3		-8.789	89.852	1.00 24.65
MOTA	5929	CB VAL B 3		-7.599	88.958	1.00 23.49
ATOM	5930	CG1 VAL B 3	68.924	-7.770	87.563	1.00 21.01
ATOM	5931	CG2 VAL B 3	69.049	-6 <i>:</i> 293	89.587	1.00 23.08
	5932	C VAL B 3				
ATOM	•			-10.083	89.144	1.00 23.19
MOTA	5933	O VAL B 3	68.691	-10.749	88.542	1.00 23.06
ATOM	5934	N LYS B 3	55 70.810	-10.436	89.216	1.00 27.15
ATOM	5935	CA LYS B 3		-11.668	88.594	1.00 29.18
MOTA	5936	CB LYS B 3		-11.758	88.704	1.00 28.61
ATOM	5937	CG LYS B 36	5 73.554	-10.617	88.030	1.00 30.27
ATOM	5938	CD LYS B 36	5 75.074	-10.768	88.154	1.00 32.58
	5939	CE LYS B 3		-9.587	87.516	1.00 29.13
ATOM						
MOTA	5940	NZ LYS B 36		-9.689	87.606	1.00 35.17
MOTA	5941	C - LYS B 36	5. 70.666	-12.879	89.276	1.00 25.30
ATOM	5942	O LYS B 36		-13.837	88613	1.00 26.81
			£ 70 EE0	12 021		1.00 26.10
MOTA	5943	N ASP B 36		-12.831	90.604	
ATOM	5944	CA ASP B 36		-13.938	91.347	1.00 28.29
ATOM	5945	CB ASP B 36	6 70.105	-13.731	92.859	1.00 29.44
ATOM	5946	CG ASP B 36		-13.669	93.311	1.00 32.95
ATOM	5947			-14.099	92.551	1.00 26.37
MOTA	5948	OD2 ASP B 36		-13.216	94.442	1.00 35.26
ATOM	5949	C ASP B 36		-14.110	90.986	1.00 28.61
					90.869	1.00 27.00
MOTA	5950			-15.231		
ATOM	5951	N THR B 36		-13.002	90.801	1.00 28.63
MOTA	5952	CA THR B 36	7 66.365	-13.080	90.438	1.00 27.35
ATOM	5953	CB THR B 36		-11.683	90.359	1.00_27.63
					91.656	
MOTA	5954			-11.068		1.00 28.12
MOTA	5955	CG2 THR B 36		-11.786	89.890	1.00 22.94
MOTA	5956	C THR B 36		-13.782	89.094	1.00 25.46
				-14.693	88.964	1.00 24.48
ATOM	5957					
ATOM	5958	N LEU B 36		-13.361	88.092	1.00 23.33
ATOM	5959	CA LEU B 36	66.857	-13.990	86.785	1.00 28.99
	5960	CB LEU B 36		-13.256	85.759	1.00 27.67
ATOM					85.046	1.00.29.47
ATOM ·	5961	CG LEU B 36		-12.070		1 00 55
ATOM	5962	CD1 LEU B 36		-12.607	84.195	1.00 32.45
ATOM	5963	CD2 LEU B 36	66.546	-11.027	86.043	1.00 19.43
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				_					
ATOM	596			B 368		2 -15.45	4 86.888	1 00	32.40
ATOM	596	5 0	LEU	B 368		-16.30		1.00	32.40
ATOM	596			B 369		3 -10.30	9 00.1/9	1.00	31.80
					08.212	2 -15.73			33.59
MOTA	596			B 369	68.656	-17.10	1 88.003	1.00	39.68
ATOM	5968	B CI	GLU	B 369	69.736	-17.14	1 89.082		42.61
ATOM	5969	9 CC		B 369	71 122	-17.13	. 00.002		44.01
ATOM	5970				71.133	-17.13	88.537		50.65
				B 369	/1.469	-18.44	3 87.842	1.00	55.81
MOTA	5973	L OF	El GLU	B 369	72.589	-18.56	1 87.299		57.42
ATOM	5972	OE	2 GLU	B 369	70.610	-19.35	87.841		58.37
MOTA	5973	3 C	GLU	B 369	67 470	-17.95			
	5974		CT	5 369	07.479	~17.954	4 88.442	1.00	34.94
MOTA			ا ټيلې	B 369	67.190	-18.974	87.827	1.00	32.71
ATOM	5975	N	LYS	B 370	66.805	-17.543	L 89.512	1 00	34.92
ATOM	5976	CA		B 370		-18.29		1.00	
ATOM	5977			B 370					35.12
•						~17.679		1.00	37.39
MOTA	5978	CG		370	65.879	-17.916	92.532	1.00	44.70
ATOM	5979	CD	LYS 1	370		-16.741			48.10
ATOM	5980	CE							
					03.336	-15.537	93.346	1.00	47.82
MOTA	5981				. 66.804	-14.387	93.786	1.00	45.41
ATOM	5982	С	LYS I	3 370	64.581	-18.375	88.930		33.21
ATOM	5983	0	LYS F	3 370	63 937	-19.409			
ATOM	5984	N	ALA E		. 64 300	-13.403	88.773		29.52
					04.390	-17.288	88.191	1.00	31.62
MOTA	5985	CA	ALA E		63.368	-17.274	87.153	1.00	37.19
ATOM	5986	CB	ALA E	371	63.392	-15.938	86.403		35.65
MOTA	5987	С	ALA E		63 572	-18.431			
ATOM	5988	ō	ALA E						37.79
						-19.137		1.00	34.46
ATOM	5989	N	LYS E		54.810	-18.644	85.759	1.00	40.10
MOTA	5990	CA	LYS B	372	65.147	-19.698	84.792	1 00	40.46
ATOM	5991	С	LYS B	372		-21.066			
ATOM	5992	ō	LYS B						43.15
					04.757	-22:053	84.591		43.57
MOTA	5993	CB	LYS B		66.654	-19.694	84.517	1.00	40.51
ATOM	5994	CG	LYS B	372	67.029	-18.925	83.248	1.00	
ATOM	5995	CD	LYS B	372	68 352	-19.390			
ATOM	5996	CE	LYS B		, 00.532	-13.350	82.635	1.00	20.00
						-20.907	82.706	1.00	20.00
ATOM	5997	NZ	LYS B		69.814	-21:354	82.116	1.00	20.00
ATOM	5998	N	ALA B	373	64.412	-21.159	86.624	1.00	
ATOM	5999	CA	ALA B	373	64 014	-22.425			
ATOM	6000	CB	ALA B		64.014	-22.425	87.239	1.00	
					64.762	-22.639	88.546	1.00	48.25
ATOM	6001	С	ALA B	373	62.515	-22.443	87.494	1.00 9	53.38
MOTA	6002	0	ALA B	373		-23.313		1.00	
ATOM	6003	OXT		373		-21.589			
HETATM		ZN	ZN C				88.269	1.00	
				1	49.660	9.211	109.302	1.00 3	32.54
HETATM		01	TSA D	2	47.669	8.189	109.464	1.00 2	28.76
HETATM	2993	C2	TSA D	2	49.952	6.981	108.340	1.00 2	
HETATM	2994	03	TSA D	2	52.458		101.667		
HETATM		N1	TSA D	2				1.00 3	
					47.800	7.789		1.00 3	
HETATM		N2	TSA D	2 .	53.013	-1.329	101.259	1.00 3	30.57
HETATM	2997	Cl	TSA D	2	51.859	2.799	101.610	1.00 2	
HETATM	2998	C2	TSA D	2	50.907		101.666	1.00 2	
HETATM		C3	TSA D	2		0.103	101.555		
				ź	51.241		101:551	1.00 2	
HETATM		C4	TSA D	2	52.626	0.026	101.366	1.00 2	13.11
HETATM		C5	TSA D	2	53.589	1.080	101.303	1.00 2	
HETATM	3002	C6	TSA D	2	53.218		101.418	1 00 2	.5.02
HETATM	3003	C7						1.00 2	
			TSA D	2	51.572		101.734	1.00 3	
HETATM	3004	C8	TSA D	2	50.108	4.726	101.996	1.00 2	9 - 05
HETATM	3005	C9	TSA D	2	50.052		103.338	1.00 2	
HETATM			TSA D	2	49.060				
							104.279	1.00 2	J.99
HETATM			TSA D	2	49.315	6.155	105.504	1.00 3	2.05
HETATM	3008		TSA D	2	48.515	6.184	106.595	1.00 2	
HETATM .	3009	C13	TSA D	2	48.855		107.756	1.00 2	
HETATM			TSA D	2					
					49.680		100.864	1.00 3	
HETATM :			TSA D	2	47.776	4.545	104.132	1.00 3	0.60
HETATM .		C17 -	TSA D	2	54.438	-1.703	101.139	1.00 2	
HETATM :		C16	TSA D	2	52.044	-2.416		1.00 2	
HETATM			ZN E	ĭ	_				
					52.949	1.842	85.681	1.00 2	
HETATM (			TSA F	2	50.964	0.911	85.428	1.00 2	
HETATM	6006	O2 '	TSA F	2	51.255		86.654	1.00 3	

	HETATM 6007 O3 TSA	F 2		51.56	9 6.51	2 02 21		
	HETATM 6008 N1 TSA	_		50.34				27.89
	HETATM 6009 N2 TSA			47.06		_		27.23
	HETATM 6010 C1 TSA			49.44				16.24
	HETATM 6011 C2 TSA			48.03				27.18
	HETATM 6012 C3 TSA			47.227				25.98
	HETATM 6013 C4 TSA			47.837				24.59
	HETATM 6014 C5 TSA							25.75
	HETATM 6015 C6 TSA	_		49.274				26.53
	HETATM 6016 C7 TSA			50.041				28.36
	HETATM 6017 C8 TSA			50.349				25.27
	HETATM 6018 C9 TSA F		•	49.716				24.18
	HETATM 6019 C10 TSA F			50.134				27.20
	HETATM 6020 C11 TSA F			49.419		•		30.21
	HETATM 6021 C12 TSA F	2		50.118				27.18
				49.762				23.47
				50.529				28.28
				50.208			1.00	28.83
				48.013			1.00	26.16
		2		47.699			1.00	27.37
	HETATM 6026 C16 TSA F			45.610			1.00	25.36
	HETATM 6027 OH2 WAT G			61.391		88.062	1.00	12.93
	HETATM 6028 OH2 WAT G			55.595			1.00	7.53
	HETATM 6029 OH2 WAT G	_		58.656	12.731	106.749	1:00	12.33
	HETATM 6030 OH2 WAT G			46.347			1.00	14.54
	HETATM 6031 OH2 WAT G	5		45.523			1.00	11.14
	HETATM 6032 OH2 WAT G	6		24.466				22.41
	HETATM 6033 OH2 WAT G	7			-17.745		1.00	21.99
		8			-15.640		1.00	26.67
		9			-14.901			23.94
		10		57.540	-7.620		1.00	26.96
		11		59.414	-2.497	84.029	1.00	22.51
		12		31.671	18.074	114.616		32.15
		13		62.335		.117.140		19.47
		14		45.565	9:469		1.00	18.81
	HETATM 6041 OH2 WAT G	15		43.311	8.237		1.00	26.11
	HETATM 6043 OH2 WAT G	16		46.628		104.423		24.28
	HETATM 6044 OH2 WAT G	17		40.672	2.507			18.30
	HETATM 6045 OH2 WAT G	18		61.830	10.923			22.27
	HETATM 6046 OH2 WAT G	19 20		57.813		108.580		24.68
	HETATM 6047 OH2 WAT G	21		48.885	5.660			30.00
	HETATM 6048 OH2 WAT G	22		36.382	-8.352	88.841	1.00	17.32
	HETATM 6049 OH2 WAT G	23		39.316	-10.091	86.422	1.00	27.38
	HETATM 6050 OH2 WAT G	24		54.802	-3.446	90.346	1.00	
	HETATM 6051 OH2 WAT G	24 25 ·		49.292		140.537	1.00	34.17
	HETATM 6052 OH2 WAT G	26		56.747	8.830	60.744	1.00	40.67
	HETATM 6053 OH2 WAT G	27		41.952	9.79	100.118	1.00	
	HETATM 6054 OH2 WAT G	28		31.268		106.695	1.00	
	HETATM 6055 OH2 WAT G	29		68.342 72.651	17.79:	111.076	1.00	30.93
	HETATM 6056 OH2 WAT G	30		39.287	-6.985	94.845	1.00	29.34
	HETATM 6057 OH2 WAT G	31		61.221	9.257	85.623	1.00	
	HETATM 6058 OH2 WAT G	32			14.462	87.256	1.00 2	
	HETATM 6059 OH2 WAT G	33		38.167 64.657			1.00	
	HETATM 6060 CH2 WAT G	34			-2.682	96.225	1.00 1	
	HETATM 6061 OH2 WAT G	35		44.059	-2.698	99.805	1.00 3	
	HETATM 6062 OH2 WAT G	36		38.480 57.899	4.763	93.051	1.00 2	
	HETATM 6063 OH2 WAT G	37				112.976	1.00.2	
	HETATM 6064 OH2 WAT G	38		57.092	3.145	93.309	1.00 2	
		38 39		52.194		118.878	1.00 3	
		40		69.400		123.379	1.00 3	
		41		24.024	6.540	79.852	1.00 3	
		41 42		46.657 -		89.402	1.00 2	
٠		42 43		24.976	13.489		1.00_4	
				46.533	-4.511	94.759	1.00 2	
		44		51.448	13.833	86.306	1.00 2	
		45 46		70.578	4.183	105.248	1.00 4	
	minin, our one wat G	46		53.938	-9.936	110.021	1.00 3	<b>8.</b> 97

									60 00F	1 00	
HETATM	6073	OH2	TAK	G	47		38.458	-0.443	63.035	1.00	28.35
HETATM	6074	OH2	WAT	G	48		64.786	7.930	107.466	1.00	34.46
HEIMIN	6075		_		49		50.823		114.809	1 00	40.51
HETATM	60/3		WAT								
HETATM	6076	OH2	WAT	G	50		33.963	-10.352	68.080		39.11
HETATM	6077	OH2	WAT	G	51		71.328	-14.321	86.007	1.00	33.30
							63.272	10.210	79.836	1 00	35.75
HETATM	60.18	OHZ	WAT	G							
HETATM	6079	OH2	VAT	G	53		59.263	-12.096	94.306	1.00	29.57
HETATM	6080	0112	WAT	C	54		46.041	10.641	76.561	1.00	27.97
HETAIM	0000								89.775		24.25
HETATM	6081		WAT	G	55			-13.620			
HETATM	6082	OH2	- WAT	G	56		76.600	0.622	89.097	1.00	29.19
HETATM	6093		WAT	G	57		53.555	6.439	79.089	1.00	34.05
HETATM	0003										35.02
HETATM	6084		WAT		58		71.301	11.026			
HETATM		OH2	WAT	G	59	•	28.188	-9.956	81.594	1.00	33.21
				G	60		53.084	20.992	98.483	1.00	27.64
HETATM											30.30
HETATM	6087	OH2	WAT	G	61		59.484	8.630			
.HETATM	6088	OH2	WAT	G	62		26.195	-3.809	95.805	1.00	33.04
HETATM		OH2	WAT	G	63		26.095	-0.121	89.620	1.00	37.39
HEIMIN	5003						47.100		109.711		20.88
HETATM	6090		WAT		64						
HETATM	6091	OH2	WAT	G	65		23.273	0.731	92.275	1.00	30.38
HETATM	6092	OH2	WAT	G	66	,	45.340	-24.751	72.694	1.00	37.51
HEIMIN	6002				67		33.754		111.676		34.63
HETATM			TAW								
HETATM	6094	OH2	TAW	G	68		52.831		126.276		47.11
HETATM				G	69		50.218	16.953	111.099	1.00	26.24
		-	WAT		70		44.791	5.844		1 00	24.95
HETATM	6096										29.48
HETATM	6097	OH2	WAT	G	71	•	49.517	-18.731			
HETATM	6098	OH2	WAT	G	72		76.379	10.131	116.550	1.00	48.70
			MAT		73	•	30.214	-8.086	87.873	1.00	46.35
HETATM								12.061			30.80
HETATM			TAW		74		45.320				
HETATM	6101	OH2	WAT	G	75		72.881	5.360	86.249	1.00	29.04
HETATM		OH2	WAT	G	76		59.674	-23.046	87.252	1.00	41.96
						•	40.619		100.345		26.45
HETATM			TAW		77						
HETATM	6104	OH2	WAT	G	78		41.666	-19.477			36.27
HETATM		OH2	WAT	G	79		46.408	-6.539	92.717	1.00	25.78
				G	80			-12:230	81.646	1.00	28.34
HETATM								-12.230	121 061		
HETATM	6107	OH2	WAT	G	81		28.268		121.961		41.15
HETATM	6108	OH2	WAT	G	82		68.843	3.154	71.986	1.00	32.34
				G	83		52 125	-11.158	85.150	1.00	24.14
HETATM	0103							-1.773	92.264		26.12
HETATM	6110	OHZ	WAT		84		75.374	-1.773	32.204		
HETATM	6111	OH2	WAT	G	85		46.957	12.230	142.271		37.07
HETATM		OH2	MAT	G	86		63.789	9.551	64.329	1.00	55.58
					87		60.672	21.185		1.00	58.55
HETATM		OH2									31.10
HETATM	6114	OH2	TAW	G	88		56.547	9.505			
HETATM	6115	OH2	WAT	G	89		26.366	-0.876	92.250	1.00	29.70
			WAT		90		67.604	-16.583	80.808	1.00	32.85
HETATM					_			1.899			42.95
HETATM	6117		WAT		91		23.910				
HET/ TM	6118	OH2	WAT	G	92		50.032	4.106	117.380	1.00	30.05
HET: IM	6119	OH2	WAT	G	93		26.774	-9.492	83.952	1.00	43.59
					94		42.714		113.787	1.00	40.17
HETA.M	0120	OH2	TAW								47.82
HETATM	6121	OH2	WAT		95		57.966		134.170		
HETATM		OH2	WAT	G	96		54.478	-3.550	119.086		36.62
HETATM			WAT		97		53.065	11.696	101.718	1.00	41.62
											45.98
HETATM	6124		WAT		98			-23.645			
HETATM	6125	OH2	WAT	G	99		54.855	-9.614	121.975	1.00	34.57
HETATM		OHO	WAT	G	100		57.408	-3.352	57.145	1.00	42.14
HEINIM	0120	0112	WAT	č	101		63.590	30 323	123.667		33.87
			UVAT	J	T 0 T		40.00	20.333	72.392		30.23
	6127	Unz		_			48.129			1.00	20.42
		OH2	WAT	G	102.			-23.143			
HETATM	6128	OH2	WAT	G G	102.		62.834	6.913			52.01
HETATM HETATM	6128 6129	OH2 OH2	WAT WAT	G	103		62.834	6.913	76.094	1.00	52.01
HETATM HETATM HETATM	6128 6129 6130	OH2 OH2 OH2	WAT WAT TAW	G G	103 104		62.834 34.566	6.913 6.529	76.094 73.089	1.00	52.01 36.29
HETATM HETATM	6128 6129 6130	OH2 OH2 OH2 OH2	TAW TAW TAW	G G G	103 104 105		62.834 34.566 51.588	6.913 6.529 20.869	76.094 73.089 67.459	1.00 1.00 1.00	52.01 36.29 36.85
HETATM HETATM HETATM	6128 6129 6130 6131	OH2 OH2 OH2 OH2	TAW TAW TAW	G G G	103 104 105		62.834 34.566	6.913 6.529	76.094 73.089 67.459 129.379	1.00 1.00 1.00	52.01 36.29 36.85 42.87
HETATM HETATM HETATM HETATM	6128 6129 6130 6131 6132	OH2 OH2 OH2 OH2 OH2	WAT WAT WAT TAW	G G G	103 104 105 106		62.834 34.566 51.588 28.160	6.913 6.529 20.869 18.020	76.094 73.089 67.459 129.379	1.00 1.00 1.00 1.00	52.01 36.29 36.85 42.87 43.62
HETATM HETATM HETATM HETATM HETATM	6128 6129 6130 6131 6132 6133	OH2 OH2 OH2 OH2 OH2 OH2	WAT WAT WAT WAT WAT WAT	G G G G	103 104 105 106 107		62.834 34.566 51.588 28.160 49.082	6.913 6.529 20.869 18.020 -11.452	76.094 73.089 67.459 129.379 57.603	1.00 1.00 1.00 1.00	52.01 36.29 36.85 42.87 43.62
HETATM HETATM HETATM HETATM HETATM HETATM HETATM	6128 6129 6130 6131 6132 6133 6134	OH2 OH2 OH2 OH2 OH2 OH2	TAW TAW TAW TAW TAW	G G G G G	103 104 105 106 107		62.834 34.566 51.588 28.160 49.082 44.717	6.913 6.529 20.869 18.020 -11.452 -8.605	76.094 73.089 67.459 129.379 57.603 93.281	1.00 1.00 1.00 1.00 1.00	52.01 36.29 36.85 42.87 43.62 41.95
HETATM HETATM HETATM HETATM HETATM HETATM HETATM	6128 6129 6130 6131 6132 6133 6134	OH2 OH2 OH2 OH2 OH2 OH2 OH2	TAW TAW TAW TAW TAW TAW	6 6 6 6 6 6	103 104 105 106 107 108		62.834 34.566 51.588 28.160 49.082 44.717 67.088	6.913 6.529 20.869 18.020 -11.452 -8.605	76.094 73.089 67.459 129.379 57.603 93.281 94.019	1.00 1.00 1.00 1.00 1.00	52.01 36.29 36.85 42.87 43.62 41.95 35.71
HETATM HETATM HETATM HETATM HETATM HETATM HETATM HETATM	6128 6129 6130 6131 6132 6133 6134 6135	OH2 OH2 OH2 OH2 OH2 OH2 OH2	TAW TAW TAW TAW TAW TAW	6 6 6 6 6 6	103 104 105 106 107 108		62.834 34.566 51.588 28.160 49.082 44.717	6.913 6.529 20.869 18.020 -11.452 -8.605 -11.900	76.094 73.089 67.459 129.379 57.603 93.281 94.019 100.800	1.00 1.00 1.00 1.00 1.00 1.00	52.01 36.29 36.85 42.87 43.62 41.95 35.71 35.40
HETATM HETATM HETATM HETATM HETATM HETATM HETATM HETATM HETATM	6128 6129 6130 6131 6132 6133 6134 6135 6136	OH2 OH2 OH2 OH2 OH2 OH2 OH2 OH2	TAW TAW TAW TAW TAW TAW TAW	9999999	103 104 105 106 107 108 109		62.834 34.566 51.588 28.160 49.082 44.717 67.088 49.561	6.913 6.529 20.869 18.020 -11.452 -8.605 -11.900	76.094 73.089 67.459 129.379 57.603 93.281 94.019 100.800	1.00 1.00 1.00 1.00 1.00 1.00	52.01 36.29 36.85 42.87 43.62 41.95 35.71 35.40
HETATM	6128 6129 6130 6131 6132 6133 6134 6135 6136	OH2 OH2 OH2 OH2 OH2 OH2 OH2 OH2 OH2	TAW TAW TAW TAW TAW TAW TAW	66666666	103 104 105 106 107 108 109		62.834 34.566 51.588 28.160 49.082 44.717 67.088 49.561 75.853	6.913 6.529 20.869 18.020 -11.452 -8.605 -11.900 22.763	76.094 73.089 67.459 129.379 57.603 93.281 94.019 100.800	1.00 1.00 1.00 1.00 1.00 1.00 1.00	52.01 36.29 36.85 42.87 43.62 41.95 35.71 35.40 56.20
HETATM HETATM HETATM HETATM HETATM HETATM HETATM HETATM HETATM	6128 6129 6130 6131 6132 6133 6134 6135 6136	OH2 OH2 OH2 OH2 OH2 OH2 OH2 OH2 OH2	TAW TAW TAW TAW TAW TAW TAW	66666666	103 104 105 106 107 108 109		62.834 34.566 51.588 28.160 49.082 44.717 67.088 49.561	6.913 6.529 20.869 18.020 -11.452 -8.605 -11.900 22.763	76.094 73.089 67.459 129.379 57.603 93.281 94.019 100.800	1.00 1.00 1.00 1.00 1.00 1.00 1.00	52.01 36.29 36.85 42.87 43.62 41.95 35.71 35.40

#### 210/263 Figure 18-94

	•
HETATM 6139 OH2 WAT G 113	33.114 1.764 67.443 1.00 37.01
HETATM 6140 OH2 WAT G 114	42.618 -4.357 102.345 1.00 39.18
HETATM 6141 OH2 WAT G 115	53.605 -10.816 66.281 1.00 31.62
HETATM 6142 OH2 WAT G 116	
HETATM 6143 OH2 WAT G 117	50 000
HETATM 6144 OH2 WAT G 118	
HETATM 6145 OH2 WAT G 119	20.00
HETATM 6146 OH2 WAT G 120	
HETATM 6147 OH2 WAT G 121	2.00 40.00
HETATM 6148 OH2 WAT G 122	
HETATM 6149 OH2 WAT G 123	
HETATM 6150 OH2 WAT G 124	41.381 -23.534 70.872 1.00 38.10
HETATM 6151 OH2 WAT G 125	31.999 1.992 73.813 1.00 33.97
HETATM 6152 OH2 WAT G 125	55.761 10.285 101.654 1.00 47.66
HETATM 6152 OH2 WAT G 126	30.596 12.964 133.642 1.00 37.98
	59.611 5.347 136.114 1.00 46.39
	24.190 12.220 124.679 1.00 30.77
HETATM 6155 OH2 WAT G 129	70.078 4.455 86.283 1.00 36.11
HETATM 6156 OH2 WAT G 130	57.882 -4.314 125.597 1.00 41.40
HETATM 6157 OH2 WAT G 131	45.838 -20.690 65.884 1.00 35.98
HETATM 6158 OH2 WAT G 132	47.574 3.186 79.027 1.00 36.67
HETATM 6159 OH2 WAT G 133	46.856 -18.901 62.295 1.00 45.40
HETATM 6160 OH2 WAT G 134	40.164 5.047 95.358 1.00 31 39
HETATM 6161 OH2 WAT G 135	27.268 -0.405 122.461 1.00 38.16
HETATM 6162 OH2 WAT G 136	54.200 -20.155 66.212 1.00 37 55
HETATM 6163 OH2 WAT G 137	45.435 -10.534 103.626 1.00 37.96
HETATM 6164 OH2 WAT G 138	31.633 25.030 106.499 1.00 43.94
HETATM 6165 OH2 WAT G 139	79.029 -7.518 93.606 1.00 40 55
HETATM 6166 OH2 WAT G 140	68.597 20.711 111.685 1.00 33.25
HETATM 6167 OH2 WAT G 141	64.263 8.524 113.832 1.00 40 63
HETATM 6168 OH2 WAT G 143	49.387 ~24.485 70.152 1.00 34.07
HETATM 6169 OH2 WAT G 144	49.387 -24.485 70.152 1.00 34.07 23.383 -3.854 83.604 1.00 32.22
HETATM 6170 OH2 WAT G 145	42.360 -0.710 61.686 1.00.35 04
HETATM 6171 OH2 WAT G 146	34.421 -3.304 65.685 1.00 35.42
HETATM 6172 OH2 WAT G 147	31.506 3.409 89.579 1.00 39.86
HETATM 6173 OH2 WAT G 148	34.963 10.688 91.806 1.00 31.12
HETATM 6174 OH2 WAT G 149	54.859 -15.085 96.769 1.00 46.65
HETATM 6175 OH2 WAT G 150	34.695 2.391 131.273 1.00 39.22
HETATM 6176 OH2 WAT G 151	40.348 1.395 61.905 1.00 34.09
HETATM 6177 OH2 WAT G 152	66.912 17.666 127.489 1.00 45.19
HETATM 6178 OH2 WAT G 153	31.096 19.900 103.232 1.00 43.45
HETATM 6179 OH2 WAT G 154	28.074 -4.222 70.175 1.00 28.86
HETATM 6180 OH2 WAT G 155	63.586 -1.894 99.003 1.00 41.15
HETATM 6181 OH2 WAT G 156	54.145 -22.222 88.415 1.00 40.92
HETATM 6182 OH2 WAT G 157	62.443 13.765 89.547 1.00 33.69
HETATM 6183 OH2 WAT G 158	58.832 9.798 101.311 1.00 31.00
HETATM 6184 OH2 WAT G 159	37.701 -5.528 119.322 1 00 45.00
HETATM 6185 OH2 WAT G 160	43.599 13.442 131.274 1 00 38.43
	23.540 -1.137 96.111 1 00 51.83
HETATM 6187 OH2 WAT G 162	59.915 -4.318 110.873 1.00 41.92
HETATM 6188 OH2 WAT G 163	51.265 -8.264 60.546 1.00 31.25.
HETATM 6189 OH2 WAT G 164	58.109 7.024 98.294 1.00 46.30
HETATM 6190 OH2 WAT G 165	46.553 18.195 74.179 1.00 37.53
HETATM 6191 OH2 WAT G 166	55.706 -21.025 92.515 1.00 43.91
HETATM 6192 OH2 WAT G 167	67.146 -1.958 109.704 1.00 43.13
HETATM 6193 OH2 WAT G 168	47.445 -3.047 134.746 1.00 27.99
HETATM 6194 OH2 WAT G 169	65.193 5.304 63.562 1.00 36.05
HETATM 6195 OH2 WAT G 170	36.176 8.979 102.024 1.00 39.63
HETATM 6196 OH2 WAT G 171	70.527 5.797 70.886 1.00 44.69
HETATM 6197 OH2 WAT G 172	
HETATM 6198 OH2 WAT G 173	
HETATM 6199 OH2 WAT G 174	
HETATM 6200 OH2 WAT G 175	
HETATM 6201 OH2 WAT G 176	
HETATM 6202 OH2 WAT G 177	
HETATM 6203 OH2 WAT G 178	
HETATM 6204 OH2 WAT G 179	
miture of of Our Mal G 1/A	52.811 11.799 98.957 1.00 36.09

HETATM	6205	OH2	WAT G	180	38.58	9 18.249	88.356	1.00 36.19
HETATM	6206	OH2	WAT G	181	43.73	4 -15.681	61.135	1.00 34.24
HETATM		OH2	WAT G	182	42.28	3 15.251	91.437	1.00 37.96
HETATM		OH2				1 -11.129		1.00 45.78
HETATM			WAT G			1 -19.367		1.00 36.55
					56.88		.95.969	1.00 39.12
HETATM		OH2						1.00 33.12
HETATM			WAT G		26.35		125.052	
HETATM		OH2			24.63	1 20.230	122.650	1.00 45.67
HETATM	6213	OH2	WAT G	.188		6 4.964		1.00 42.16
HETATM	6214	OH2	WAT G	189	55.01	7 14.964	62.948	1.00 50.18
HETATM	6215	OH2	WAT G	190	33.37	1 13.710	105.640	1.00 37.04
HETATM		OH2	WAT G	191	44.46	6 -10.386	91.144	1.00 36.62
HETATM			WAT G		28.43		121.285	1.00 38.19
HETATM	6218		WAT G		29.78	6 24.957	122.112	1.00 42.05
HETATM	6219		WAT G		28.85	2 3.461		1.00 48.35
			WAT G		41 69	1 11.318		1.00 32.60
HETATM			WAT G		25 01	1 11.318 2 -10.229 2 -23.250	111 631	1.00 47.70
HETATM					40.01	2 -10.223	76 630	1.00 48.86
HETATM		OH2				2 -23.250	76.629	
HETATM			WAT G				121.410	1.00 43.09
HETATM			WAT G			4 -20.514		1.00 46.80
HETATM	6225	OH2		200		3 -13.079		1.00 44.89
HETATM	6226	OH2	WAT G	201	59.98	2 24.381	103.984	1.00 40.63
HETATM		OH2	WAT G	202		412.771	74.705	1.00 35.13
HETATM		OH2	WAT G	203	28.70	8 9.2 <b>1</b> 1	79.238	1.00 33.53
HETATM			WAT G		53.25		122.243	1.00 48.49
HETATM			WAT G		50.70			1.00 41.26
HETATM	6231		WAT G		50.00			1.00 39.15
		OH2				8 -16.236		1.00 29.70
HETATM			WAT G		24.39		111.635	1.00 50.82
HETATM							114.289	1.00 44.49
HETATM			WAT G		53.38			1.00 31.97
HETATM			WAT G		60.12			
HETATM			WAT G		23.40		111.744	1.00 44.97
HETATM			WAT G		46.21		76.878	1.00 59.14
HETATM	6238		WAT G		29.75		97.109	1.00 41.78
HETATM	6239	OH2	WAT G	214	46.82			1.00 41.02
HETATM	6240	OH2	WAT G	215	59.14		124.775	1.00 38.42
HETATM		OH2	WAT G	216	42.67	4 14.088	66.037	1.00 32.50
HETATM		OH2	WAT G	217	55.00			1.00 56.50
HETATM		OH2	WAT G	218	63.36	1 -8.209	109.653	1.00 49.66
HETATM		OH2	WAT G	219	66.58	3 ~8.146	94.671	1.00 50.91
HETATM			WAT G			72.583		1.00 36.99
HETATM			WAT G		24.47			1.00 47.24
HETATM			WAT G		76.91			1.00 50.43
			WAT G		32.78	9 0 651	129.136	1.00 42.47
HETATM			WAT G			1 -16.880		1.00 46.69
HETATM			WAT G		78.56			1.00 43.17
HETATM	6250							1.00 35.84
HETATM			WAT G		45.68			
HETATM	6252			227	38.26	3 15.236	84.711	1.00 42.39
HETATM	6253		WAT G		38.93		108.488	1.00 52.23
HETATM	6254		WAT G		33.75	5 14.939	70.228	1.00 46.56
HETATM	6255	OH2	WAT G	230	51.52	1 34.184	100.859	1.00 52.96
HETATM	6256		WAT G		34.14	0.565	63.039	1.00 31.02
HETATM	6257	он2	WAT G	232	37.27	7 13.977	81.662	1.00 39.83
HETATM'	6258		WAT G		57.30			1.00 28.55
HETATM	6259		WAT G		31.71			1.00 53.16
WEIWIN	6760		WAT G		60.62			1.00 51.02
HETATM			WAT G		44.35		94.916	1.00 57.50
HETATM	0201		WAT G				70.111	1.00 38.65
HETATM	0202				68.45		79.253	1.00 44.95
HETATM	6263		WAT G		27.83		81.087	
HETATM	6264		WAT G		. 54.93			1.00 42.95
HETATM	6265		WAT G			2 -15.271	71.579	1.00 45.78
HETATM	6266		WAT G		35.96		61.973	1.00 38.91
HETATM	6267		WAT G		29.68			1.00 44.85
HETATM	6268	он2	WAT G	243	49.53	4 -10.150	113.501	1.00 38.32
HETATM	6269	OH2	WAT G	244	57.25	2 9.773	96.696	1.00 48.83
HETATM	6270		WAT G		62.31		80.972	1.00 38.54
UPIVIU	J			_			•	

HETAT	M 627	L OH2 WAT G 246		50.248	~5.552	2 102.815	1.00 43.23
HETAT	M 6272	OH2 WAT G 247		47.966			
HETAT	M 6273	OH2 WAT G 248		62.507		5 108.414	1.00 35.30
	M 6274			53.971	19.763		1.00 47.49
	M 6275			38.406	9.828		1.00 33.71
	M 6276			35.304	-6 170	_	
				35.304	-6.179		1.00 36.23
	M 6277			39.218	-12.667		1.00 36.17
	M 6278			56.350		-	1.00 46.38
	M 6279			69.850		122.119	1.00 55.07
	M 6280			75.703		128.600	1.00 30.64
HETAT	м 6281	OH2 WAT G 256	-	32.019	-12.973	113.965	1.00 34.48
HETAT	M 6282	OH2 WAT G 257		54.081	3.421	56.994	1.00 39.11
HETAT	M.6283	OH2 WAT G 258		32.801		91.078	1.00 35.72
HETAT	M 6284	OH2 WAT G 259		45.040 39.815	0.301	95.449	1.00 36.57
HETAT	M 6285	OH2 WAT G 260		39.815	21.460	128.855	1.00 40.10
HETAT	M 6286	OH2 WAT G 261		28.763	10.408	93.790	1.00 44.39
	M 6287			49.668	-12.050	93.790 60.539	1.00 50.89
	M 6288			64 252	20 015	117 400	1.00 62.67
	M 6289			75.183 46.289	13 021	128.124	1.00 50.42
	M 6290	OH2 WAT G 265	•	16 220	5 926	52.485	1.00 30.42
	M 6291	OH2 WAT G 266		68.708	13.973	70.958	
	M 6292			71.504	13.3/3		1.00 37.90
		OH2 WAT G 268		36.309		130.029	1.00 38.78
	M 6293					130.364	1.00 42.92
	4 6294	OH2 WAT G 269		65.973	12.195		1.00 51.68
	4 6295	OH2 WAT G 270		71.952			1.00 37.70
	4 6296	OH2 WAT G 271.		44.433	~17.578		1.00 49.33
	4 6297	OH2 WAT G 272		26.917 63.380		•	1.00 38.07
	4 6298	OH2 WAT G 273		63.380	-5.416	126.550	
HETATI		OH2 .WAT G 274		63.360	-5.356	95.641	1.00 37.54
	4 6300	OH2 WAT G 275		63.360 65.947	-13.015	97.485	1.00 37.42
HETATI		OH2 WAT G 276		26.406	25.831	117.328	1.00 48.37
HETATI		OH2 WAT G 277		41.893	-10.251	98.201	1.00 46.36
HETATI	6303	OH2 WAT G 278		30.343	-6.507	117.764	1.00 49.87
HETATI	1 6304	OH2 WAT G 279.		45.135	32:419	111.056	1.00 43.93
HETATI	6305	OH2 WAT G 280		50.553	-1.365	120.511	1.00 54.02
HETATI	6306	OH2 WAT G 281		60.428	13.652	105.130	1.00 31.10
HETATM	6307	OH2 WAT G 282		30.342	2.204		1.00 45.19
HETATM	6308	OH2 WAT G 283		60.358		127.736	1.00 33.17
HETATM	6309	OH2 WAT G 284		64.193	3.421	62.117	1.00 45.81
HETATM	6310	OH2 WAT G 285		45.468		105.853	1.00 48.98
HETATM	6311	OH2 WAT G 286		47.514	3.808	98.279	1.00 46.45
HETATM		OH2 WAT G 287		72.144	-6.345		1.00 40.04
HETATM		OH2 WAT G 288		54.142	-5.100		1.00 43.62
HETATM		OH2 WAT G 289		48.982	13.297	65.822	1.00 46.98
HETATM		OHZ WAT G 290		41.171	34.107		1.00 51.76
HETATM		CH2 WAT G 291		36.494	37.195		1.00 44.27
HETATM		OH2 WAT G 292		48.580	23.117	85.456	1.00 40.96
HETATM		OH2 WAT G 293		55.853	22.934		1.00 40.95
HETATM		OH2 WAT G 294		61.720	11.077		1.00 40.93
HETATM		OH2 WAT G 295				114.112	1.00 41.21
HETATM		OH2 WAT G 296				129.052	
				53.001			1.00 37.41
HETATM		OH2 WAT G 297		70.258	24.928	82.843	1.00 48.09
HETATM		OH2 WAT G 298		77.493		130.507	1.00 51.77
HETATM		OH2 WAT G 299		32.233	12.182	83.028	1.00 53.51
HETATM		OH2 WAT G 300		40666	12.878	65.747	1.00 46.49
HETATM		OH2 WAT G 301		50.977		114.597	1.00 48.51
HETATM		OH2 WAT G 302		54.236	3.817	92.196	1.00 41.15
HETATM		CH2 WAT G 303		59.527	-1.343	107.471	1.00 36.71
HETATM		OH2 WAT G 304		70.331	3.940	89.312	1.00 47.70
HETATM		CH2 WAT G 305		60.626	6.969	127.780	1.00 41.96
MTATH		OH2 WAT G 306		42.156	-0.139	133.156	1.00 32.19
HETATM		OH2 WAT G 307		58.886	16.514	99.413	1.00 53.60
HETATM		OH2 WAT G 308		67.617	-1.589	96,570	1.00 40.36
HETATM		OH2 WAT G 309			-10.936	98.849	1.00 48.80
HETATM		OH2 WAT G 310		45.576		131.914	1.00 48.99
METATM		CH2 WAT G 311		37.583	-6.243	64.257	1.00 37.06
					V. 2.2		

HETATM	1 6337	OH:	2 WA:	rc	312		66.759	16.408	94.600	1.00	45.07
HETATM	r 6338	OH:	2 WAS	rc	313		24.142	11.212	113.340	1.00	52.23
HETATM					314		69.409			-	39.88
HETATM	1 6340				315		22.064	24.858	115.328	1.00	50.23
HETATM	6341	OH	CAW S	rc	316		50.171	9.551	100.345	1.00	37.32
HETATM					317		55.104		119.497		44.78
HETATM					318		65.333	-10.105		1.00	44.21
HETATM	6344	OH2	CAW S	re	319		31.415	-2.472	128.127	1.00	41.95
HETATM		OH	י שא	r c	320		37.423	13.143	88.069		44.79
•											
HETATM	6 6346				321		43.619		96.509		54.69
HETATM	6347	OH2	CAW S	r G	322	•	68.048	14.555	126.016	1.00	42.75
HETATM	6348	OH2	LAM S		323		34.778	-2.509	130.204	1. 00	37.06
					324						
HETATM							27.972		103.841		47.34
HETATM	6350	OHZ	RAW S	. G	325		53.550	23.610	97.592	1,00	38.03
HETATM	6351	OH2	CAW S	· G	326		33.776	4.171	103.451	1.00	50.60
HETATM					327		37.862		114.870		48.34
					-						
HETATM	6353	OHZ	: WAI	' G	328		50.893	14.612	93.478	1.00	38.77
HETATM	6354	OH2	LAW :	' G	329		71.422	-20.913	86.137	1.00	47.69
HETATM			. WAI		330			-23.133	74.502		41.94
							30.310				
HETATM			raw :				41.520	7.269	60.583		54.93
HETATM	6357	OH2	LAW:	, G	332		75.879	13.737	106.089	1.00	44.65
HETATM		OH2	WAT	· G	333		51.923	9.027			41.08
HETATM			WAT				49.511	27.611	79.363		39.05
HETATM			WAT				69.385	0.852	110.192		41.42
HETATM	6361	OH2	WAT	G	336		40.952	2.479	101.880	1.00	42.50
HETATM			WAT				32.998		103.784		54.22
HETATM			WAT				54.366		136.205		52.69
HETATM	6364	OH2	WAT	'G	339		35.674	13.727	89.792	1.00	35.83
HETATM	6365	OH2	WAT	G	340		66.606	-21.361	87.138	1.00	46.26
HETATM			WAT				72.053		131.550		45.27
HETATM			WAT	_			28.072	-1.358	70.419		34.92
HETATM	6368	OH2	TAW	G	343		23.611	-3.981	76.422	1.00	52.39
HETATM	6369	OH2	WAT	G	344		53.684	2.564	122.150	1.00	58.16
			WAT				30.624		125.556		34.71
HETATM	6370										
HETATM	6371	OH2	WAT	G	346		27.870	13.838	113.997	1.00	44.91
HETATM	6372	OH2	WAT	G	347		31.903	-9.588	116.327	1.00	55.34
HETATM		OH2	WAT	G	348		71.763	15.094	63.739	1.00	48.99
							25.258				
HETATM			WAT		349				114.760		37.19
HETATM	6375	OH2	WAT	G	350		43.765	12.162	78.143	1.00	42.32
HETATM	6376	OH2	WAT	G	351		32.452	5.338	73.909	1.00	33.70
HETATM	6377	OH2	WAT	G	352		52.896	~5.770	101.894	1 00	46.40
nginin	6370										
HETATM	03/8		WAT		353		47.968	4.242			34.62
HETATM	6379	OH2	WAT	G	354		38.561	-9.302	90.596	1.00	49.80
HETATM	6380	OH2	WAT	G	355		63.791	17.454	74.354	1.00	56.40
HETATM	6381		WAT		356		41.360		133:760		50.00
UPIAIN	22.02										
HETATM	6382		TAW		357		42.467	-7.937			38.01
HETATM	6383	OH2	WAT	G	358		50.890	-0.362	116.668	1.00	39.26
HETATM		OH2	TAW	G	359		54.217	-23.881	67.865	1.00	55.18
HETATM			WAT				64.959	9.539			38.83
nelaim	0703										
HETATM	6386		WAT					-19.846	82.288		38.60
HETATM	6387	OH2	WAT	G	362		42.245	-1.140	93.572	1.00	31.47
HETATM	6388	OH2	TAW	G	363		73.552	17.770	125.885	.1.00	54.89
HEIMIN MI	6200										
HETATM	6369		WAT.				68.769		106.810		45.53
HETATM	6390	OH2	WAT	G	365		37.543	19.031	78.866	1.00	45.15
HETATM	6391	OH2	WAT	G	366		55.583	6.906	95.087	1.00	44.99
HETATM	6392		WAT				41.284	9.699	78.250		
BLININ	C3C3									-	
HETATM	0227		WAT				25.203	5.332	126.362		46.60
HETATM	6394	OH2	WAT	G	369		74.742	~5.006	95.104	1.00	47.85
HETATM	6395	OH2	TAW	G	370		70.349	19.871	69.925		51.46
*********	6306		WAT				42.93.6	20.631	94.720		38.66
HETATM	0250									_	
HETATM	6597				372			-16.114	114.141		44.01
HETATM	6398	OH2	TAW	G	373		33.863	16.838	100.275		44.66
HETATM	6399	OH2			374		21.613	12.569	86.140		43.89
BEIVIN	6400								100.583	1 00	53:53
HETATM	0400		TAW								
PETATM	6401		WAT				70.095	13.395	117.505		52.02
HETATM	6402	OH2	WAT	G	377		41.853	19.108	131.799	1.00	46.47
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HETATM 6403 OH2 WAT G 378	FR 844 44 44 44 44
	55.780 -14.986 65.487 1.00 49.09
HETATM 6404 OH2 WAT G 379	40.990 21.205 91.611 1.00 41.02
HETATM 6405 OH2 WAT G 380	48.157 1.057 116.992 1.00 44.84
HETATM 6406 OH2 WAT G 381	
	30.221 27.743 109.194 1.00 39.92
HETATM 6408 OH2 WAT G 383	49.926 -12.826 118.421 1.00 58.95
HETATM 6409 OH2 WAT G 384	42.435 -17.636 81.477 1.00 48.47
HETATM 6410 OH2 WAT G 385	
,	58.226 -25.990 71.378 1.00 48.18
HETATM 6411 OH2 WAT G 386	40.495 17.944 128.741 1.00 43.82
HETATM 6412 OH2 WAT G 387	31.943 6.301 109.475 1.00 35.53
HETATM 6413 OH2 WAT G 388	47.277 2.559 100.509 1.00 43.00
HETATM 6413 OH2 WAT G 388 HETATM 6414 OH2 WAT G 389	47.277 2.559 100.509 1.00 43.00 38.862 9.112 102.620 1.00 31.70
METATH 0414 OH2 WAI G 389	38.862 9.112 102.620 1.00 31.70
HETATM 6415 OH2 WAT G 390.	71.652 14.568 105 167 1 00 49 63
HETATM 6416 OH2 WAT G 391	68.554 -10.518 73.331 1.00 38.16
HETATM 6417 OH2 WAT G 392	70.496 -16.160 84.425 1.00 32.16
	70.496 -16.160 84.425 1.00 32.16
HETATM 6418 OH2 WAT G 393	44.698 -24.950 75.603 1.00 43.38
HETATM 6419 OH2 WAT G 394	56.172 15.369 55.027 1.00 47.44
HETATM 6420 OH2 WAT G 395	46.150 -9.441 99.999 1.00 47.98 26.892 -8.356 89.057 1.00 34.99
HETATM 6421 OH2 WAT G 396	26 002 0 354 00 057 1 00 34,36
	26.892 -8.356 89.057 1.00 34.99
HETATM 6422 OH2 WAT G 397	31.737 14.380 90.395 1.00 50.78 36.261 -13.824 62.777 1.00 50.86
HETATM 6423 OH2 WAT G 398	36.261 -13.824 62.777 1.00 50.86
HETATM 6424 OH2 WAT G 399	37.312 15.242 134.977 1.00 43.57
HETATM 6425 OH2 WAT G 400	
	33.728 13.773 126.419 1.00 57.13
HETATM 6426 OH2 WAT G 401	45.269 27.937 130.311 1.00 49.55
HETATM 6427 OH2 WAT G 402	44.887 -17.414 111.508 1.00 54.29
HETATM 6428 OH2 WAT G 403	68.928 0.455 136.711 1.00 49.90
HETATM 6429 OH2 WAT G 404	
	43.271 -21.571 64.425 1.00 48.61
HETATM 6430 OH2 WAT G 405	24.243 -4.781 108.590 1.00 51.05
HETATM 6431 OH2 WAT G 406	54.828 5.311 59.009 1.00 43.43
HETATM 6432 OH2 WAT G 407	53.460 27.992 124.076 1.00 47.83
HETATM 6433 OH2 WAT G 408	70.833 -18.390 85.386 1.00 49.26
	70.833 -18.390 85.386 1.00 49.26
HETATM 6434 OH2 WAT G 409	71.497 15.287 113.071 1.00 34.52
HETATM 6435 OH2 WAT G 410	36.407 -18.480 110.466 1.00 55.43
HETATM 6436 OH2 WAT G 411	26.220 -9.551 78.158 1.00 47.69
HETATM 6437 OH2 WAT G 412	
	52.319 26.326 82.038 1.00 42.00
HETATM 6438 OH2 WAT G 413	76.173 14.097 122.253 1.00 44.90
HETATM 6439 OH2 WAT G 414	58.379 6.335 123:024 1.00 54.61
HETATM 6440 OH2 WAT G 415	72.162 -16.705 82.719 1.00 50.63
HETATM 6441 OH2 WAT G 416	63.557 26.152 65.944 1.00 39.83
HETATM 6442 OH2 WAT G 417	30 035 23 050 100 740 7 00 75
	38.935 23.070 122.742 1.00 52.57
HETATM 6443 OH2 WAT G 418	55.256 -10.714 124.501 1.00 42.38
HETATM 6444 OH2 WAT G 419	55.443 -9.037 110.170 1.00 46.47
HETATM 6445 OH2 WAT G 420	73.873 16.578 123.288 1.00 46.54
HETATM 6446 OH2 WAT G 421	
	74.426 12.663 117.527 1.00 43.62
HETATM 6447 CH2 WAT G 422	52.374 -0.368 51.502 1.00 56.99
HETATM 6448 OH2 WAT G 423	60.339 20.215 84.713 1.00 36.27
HETATM 6449 OH2 WAT G 424	48.308 1.354 54.561 1.00 38.53
HETATM 6450 OH2 WAT G 425	
HETATM 6451 OH2 WAT G 426	33.222 -14.916 119.528 1.00 51.12
HETATM 6452 OH2 WAT G 427	47.477 3.359 112.298 1.00 46.10
HETATM 6453 OH2 WAT G 428	39.909 2.272 138.388 1.00 35.33
HETATM 6454 OH2 WAT G 429	57.000 15.272 130.300 1.00 33.33
	57.829 15.336 126.262 1.00 62.59
HETATM 6455 OH2 WAT G 430	48.917 -5.857 119.191 1.00 51.45
HETATM 6456 OH2 WAT G 431	44.139 -3.812 132.964 1.00 44.91
HETATM 6457 OH2 WAT G 432	38.885 18.594 95.398 1.00 50.23
	50.000 10.094 00.000 1,00 30.23
	52.628 -7.064 55.271 1.00 38.96
HETATM 6459 OH2 WAT G 434	60.644 -0.731 101.129 1.00 47.30
HETATM 6460 OH2 WAT G 435	64.772 5.808 71.942 1.00 50.81
HETATM 6461 OH2 WAT G 436	39.571 16.705 80.180 1.00 34.07
	120 701 0 551 65 271 3 00 47
HETATM 6462 OH2 WAT G 437	32.791 -0.551 65.371 1.00 41.40
HETATM 6463 OH2 WAT G 438	58.318 -7.989 60.087 1.00 46.94
HETATM 6464 OH2 WAT G 439	26.982 5.474 120.408 1.00 46.28
HETATM 6465 OH2 WAT G 440	72.138 1.233 90.050 1.00 50.13
HETATM 6466 OH2 WAT G 441	
	29.494 10.971 118.393 1.00 56.30
HETATM 6467 OH2 WAT G 442	69.232 5.594 113.941 1.00 58.17
HETATM 6468 OH2 WAT G 443	61.459 11.576 71.140 1.00 61.67
•	•

HETATM	6469	OH2	WAT	G	444	59.592	2.195	58.518	1.00	42.66
HETATM	6470	OH2	_WAT	G	445	47.407	6.152	111.310	1.00	45.14
HETATM	6471	OH2	WAT	G	446	36.254	18.203	99.930	1.00	44.76
HETATM	6472	OH2	WAT	G	447	49.525	32.050	116.235	1.00	47.72
HETATM	6473	OH2	WAT	G	448	21.801	-5.358	81.109	1.00	42.07
HETATM	6474	OH2	WAT	G	449	52.131	-14.007	95.380	1.00	40.76
HETATM	6475	OH2	WAT	G	450	39.712	-19.983	72.499	1.00	51.69
HETATM	6476	OH2	TAW	G	451	67.651	5.620	67.102	1.00	42.38
HETATM	6477	OH2	WAT	G	452	77.344	1.313	79.207	1.00	63.64
HETATM	6478	OH2	WAT	G	453	55.249	-29.426	86.187	1.00	44.98
HETATM	6479	OH2	WAT	G	454	64.429	-11.004	98.104	1.00	49.12
HETATM	6480	OH2	WAT	G	455	45.456	-0.814	129.510	1.00	61.60
HETATM	6481	OH2	WAT	G	456	65.066	-14.790	68.028	1.00	40.08
HETATM	6482	OH2	WAT	G	457	34.732	5.611	94.924	1.00	58.32

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#### Figure 19-1

	· Do-44	- 4 V	-	_	_	
ATOM	Residu		Y	_ Z	B Segment	ID
		45.86				AAAA
ATOM		. 46.76				AAAA
ATOM	3 0 ALA A 2	46.33				AAAA
ATOM	4 :: ALA A 2	48.28			1.00 57.26	AAAA
ATOM	5 CA ALA A 2	47.06	2 37.537	7 74.110		AAAA
ATOM	6 H LYS A 3	46.97	5 .38.628	3 71.938		AAAA
АТСИ	7 CA LYS A 3	46.72	1 39.716	71.002		AAAA
ATOM	8 CE LYS A 3	47.81				
ATOM	9 CG LYS A 3	49.22				AAAA
ATOM	10 CD LYS A 3	50.252				AAAA
ATOM	11 CE LYS A 3	51.654				AAAA
ATOM	12 NZ LYS A 3	52.643				AAAA
ATOM	13 C LYS A 3	45.393				AAAA
ATOM	14 0 LYS A 3					. YYYY
ATOM		44.894				AAAA
		44.826				AAAA
ATOM	16 CA VAL A 4	43.561				AAAA
ATOM	17 CB VAL A 4	42.543			1.00 42.26	AAAA
ATOM	18 CG1 VAL A 4	41.213		68.940		AAAA
ATOM	19 CG2 VAL A 4	42.401	41.307	71.128	1.00 42.00	AAAA
ATOM	20 C VALA 4	43.918	40.913	67.638	1.00 39.94	AAAA
MOTA	21 0 VAL A 4	44.332	42.032	67.395	1.00 40.39	AAAA
MOTA	22 n lysa 5	43.766	40.001	66.695	1.00 36.94	AAAA
ATOM	23 CA LYS À 5	44.142	40.305	65.323	1.00 34.10	AAAA
ATOM	24 CB LYS A 5	45.179			1.00 35.02	AAAA
ATOM	25 CG LYS A 5	46.424			1.00 34.07	
ATOM	26 CD LYS A 5	47.233	40.452	65.652	1.00 33.44	AAAA
MOTA	27 CE LYS A 5	48.555	40.239		1.00 32.38	aaaa
MOTA	28 NZ LYS A 5	49.372	41.460	66.222	1.00 31.26	AAAA
ATOM	29 C LYS A 5	42.997		64.333	1.00 31.38	AAAA
ATOM	30 O LYS A 5	42.053	39.523	64.466	1.00 31.38	AAAA
ATOM	31 N LEU A 6	43.090	41.142	63.326	1.00 28.60	AAAA
ATOM	32 CA LEU A 6	42.075	41.142	62.289		AAAA
ATOM	33 CB LEU A 6	41.530	42.580		1.00 26.90	AAAA
ATOM	34 CG LEU A 6	40.321			1.00 25.43	AAAA
ATOM	35 CD1 LEU A - 6		42.748	61.129	1.00 25.89	AAAA
ATOM	36 CD2 LEU A 6	40.108	44.224	60.826	1.00 25.50	AAAA
ATOM		40.550	42.032	59.828	1.00 26.64	<b>EAA</b> K
	37 C LEU A 6 38 C LEU A 6	42.818	40.701	61.049	1.00 25.74	AAAA
ATOM		43.877	41.226	60.717	1.00 24.60	AAAA
ATOM		42.282	39.704	60.357	1.00 25.90	AAAA
ATOM	40 CA ILE A 7	42.939	39.212	59.173	1.00 26.75	. AAAA
ATOM	41 CB ILE A 7	42.839	37.712	59.089	1.00 26.58	AAAA
ATOM	42 CG2 ILE A 7	43.474	37.227	57.783	1.00 27.88	AAAA
ATOM	43 CG1 ILE A ?	43.528	37.116	60.310	1.00 27.02	AAAA
ATOM	44 CD1 TLE A 7	43.507	35.640	60.350	1.00 27.46	AAAA
ATOM	45 C ILE A 7	42.339	39.814	57·. 929	1.00 26.70	AAAA
MOTA	46 0 ILE A 7	41.162	39.655	57.681	1.00 27.68	AAAA
atom	47 N GLY A 8	43.144	40.509	57.142	1.00 27.94	AA A
ATCM	48 CA GLY A 8	42.598	41.110	55.944	1.00 29.78	AA: A
ATOM	49 3 GLY A 8	43:.587	41.789	55.027	1.00 30.38	AAAA
ATOM	50 0 GLY A 8	44.785	41.765	55.264	1.00 29.39	AAAA
ATOM	51 N THRA 9	43.051	42.395	53.971	1.00 31.84	AAAA
ATCM	52 CA THR A 9	43.832	43.106	52.962	1.00 32.41	AAAA
ATCM	53 CB THR A 9	44.606	42.112	52.064	1.00 31.12	AAAA
ATOM	54 CG1 THR A 9	45.324	42.825	51.053	1.00 30.74	AAAA
ATOM	55 CG2 THR A 9	43.654	41.140	51.411	1.00 30.27	AAAA
ATOM	56 C THR A 9	42.886	43.939	52.091	1.00 30.27	
ATOM	57 3 THR A 9	41.705		51.993		AAAA
	58 N LEU A 10	43.396	43.625		1.00 33.62	AAAA
ATOM			45.009	51.485	1.00 33.20	AAAA
MOTA		42.573	45.840	50.611	1.00 33.29	AAAA
ATOM	60 CB LEU A 10	43.117	47.275	50.484	1.30 33.12	AAAA
ATCM	51 CG LEU A 10	43.142	48.245	51.566	1.00 32.95	AAAA
ATOM	62 CD1 LEU A 10	41.743	48.386	52.288	1.00 31.99	AAAA
atom	63 CD2 LEU A 10	44.126	47.734	52.675	1.00 34.71	AAAA
atcm	64 C LEU A 10	42.527	45.231	49.218	1.00 33.18	AAAA
ATCM	65 0 LEU A 10	41.876	45.768	48.328	1.00 32.52	AAAA
atom	66 N ASP A 11	43.230	44.121	49.022	1.00 33.56	AAAA,
	_•			-		

ATOM	67	CA	ASP :	A 11	43.240	43.489	47.716	1.00 34.24	AAAA
ATOM	68	CB	ASP A	A 11	44.393	42.499	47.607	1.00 35.81	AAAA
	69	CG	ASP		45.739	43.190	47.604	1.00 37.57	AAAA
MOTA						44.178	46.855	1.00 37.95	AAAA
ATOM	70		ASP A		45.890				
MOTA	71		ASP A		46.650	42.750	48.332	1.00 40.31	AAAA
- MOTA	72	С	ASP A		41.929	42.813	47.341	1.00 34.03	AAAA
MOTA	73	0	ASP A	A 11	41.629	42.652	46.150	1.00 34.80	AAAA
ATOM	74	N	TYR I	A 12	41.142	42.417	48.335	1.00 32.34	AAAA
ATOM	75	CA	TYR I	A 12	39.871	41.803	48.017	1.00 32.53	AAAA
ATOM	76	СВ	TYR A	_	39.043	41.569	49.290	1.00 31.32	AAAA
	77	CG	TYR		39.551	40.438	50.162	1.00 29.95	AAAA
MOTA					39.983	40.669	51.469	1.00 28.52	AAAA
MOTA	78		TYR A						AAAA
MOTA	79		TYR A		40.413	39.614	52.279-	1.00 28.03	
MOTA	80		TYR 1		39.568	39.128	49.688	1.00 28.47	AAAA
MOTA	81	CE2	TYR A		39.992	38.083	50.483	1.00 28.47	AAAA
ATOM	. 82	CZ	TYR A	A 12	40.408	38.330	51.775	1.00 28.43	- AAAA
ATOM	83	OH	TYR A	A 12	40.786	37.277	52.569	1.00 29.86	AAAA
ATOM	84	C	TYR 2	A 12	39.146	42.749	47.066	1.00 33.16	. AAAA
ATOM	85	ō	TYR A		38.554	42.324	46.082	1.00 33.36	AAAA
	86	N	GLY A		39.237	44.041	47.356	1.00 34.76	AAAA
ATOM			GLY A		38.594	45.065	46.546	1.00 36.60	AAAA
MOTA	87	CA					45.052	1.00 37.85	AAAA
ATOM	88	С	GLY A		38.814	44.961			
MOTA	89	0	GLY A		38.105	45.591	44.275	1.00 37.40	AAAA
MOTA	90	N	LYS A		39.799	44.171	44.647	1.00 39.55	AAAA
MOTA	91	CA	LYS A			43.981	43.231	1.00 40.66	AAAA
ATOM	92	CB	LYS A	A 14	41.605	43.977	42.995	1.00 42.26	AAAA
MOTA	93	CG	LYS A	A 14	42.300	45.309	43.239	1.00 44.54	AAAA
ATOM	94	CD	LYS A	A 14	41.820	46.445	42.304	1.00 46.32	AAAA
ATOM	95	CE	LYS A	A 14	42.033	46.158	40.810	1.00 46.64	AAAA
ATOM	96	NZ	LYS A		41.133	45.086	40.256	1.00 47.23	AAAA
ATOM	97	С	LYS A		39.499	42.675	42.707	1.00 40.35	AAAA
ATOM	98	ŏ	LYS A		39.593	42.377	41.511	1.00 39.97	AAAA
	99	N	TYR A		38.897	41.901	43.605	1.00 39.95	AAAA
MOTA				_		40.617	43.245	1.00 40.30	AAAA
MOTA	100	CA	TYR A		38.962	39.490	44.050	1.00 38.46	AAAA
ATOM	101	CB	TYR A			39.519	44.021	1.00 37.01	AAAA
MOTA	102	CG	TYR A		40.472			1.00 36.24	AAAA
MOTA	103		TYR A		41.213	39.136	45.137		AAAA
ATOM	104		TYR A		42.604	39.220	45.144	1.00 35.73	
MOTA	105		TYR A		41.163	39.976	42.902	1.00 36.84	AAAA
MOTA	106		TYR A		42.556	40.064	42.898	1.00 36.53	AAAA
MOTA	107	CZ	TYR A	A 15	43.271	39.689	44.028	1.00 36.24	AAAA
MOTA	108	OH	TYR A	A 15	44.648	39.816	44.042	1.00 36.49	AAAA
ATOM	109	С	TYR A	A 15	36.802	40.647	43.556	1.00 41.98	AAAA
ATOM	110	0	TYR A	A 15	36.288	39.786	44.280	1.00 42.59	AAAA
ATOM	111	N	ARG A	A 16	36.101	41.638	43.014	1.00 42.81	· AAAA
ATOM	112	CA	ARG A		34.670	41.753	43.257	1.00 43.47	AAAA
MOTA	113	CB	ARG A	_	34.205	43.197	43.111	1.00 45.27	AAAA
	114	CG	ARG A		35.021	44.234	43.833	1.00 48.06	AAAA
ATOM	115	CD	ARG A		34.891	44.196	45.339	1.00 49.63	AAAA
ATOM					35.632	45.322	45.905	1.00 51.65	AAAA
ATOM	116	NE	ARG A		35.382	46.602	45.622	1.00 52.71	AAAA
MOTA	117	CZ	ARG A					1.00 53.28	AAAA
MOTA	118		ARG A		34.406	46.931	44.781	1.00 53.43	
ATOM	119		ARG A		36.124	47.560	46.162	1.00 53.43	AAAA
ATOM	120	С	ARG A	1 16	33.913	40.929	42.230	1.00 42.86	AAAA
ATOM	121	0	ARG A	16	34 .455	40.541	41.193	1.00 41.83	AAAA
ATOM	122	N	TYR A	17	32.651	40.668	42.523	1.00 42.42	AAAA
ATOM	123	CA	TYR A		31.818	39.942	41.590	1.00 42.76	AAAA
ATOM	124	CB	TYR A		30.675	39.254	42.333	1.00 40.11	AAAA
	125	CG	TYR A		31.097	38.061	43.180	1.00 38.35	AAAA
ATOM			TYR A		32.169	38.148	44.071	1.00 36.15	AAAA
ATOM	126				32.519	37.069	44.874	1.00 34.76	AAAA
MOTA	127	CE1	TYR A		30.386	36.855	43.116	1.00 36.40	AAAA
ATOM	128	CD2	TYR A		30.726	35.776	43.110	1.00 35.31	AAAA
MOTA	129	CE2	TYR A					1.00 35.00	AAAA
MOTA	130	CZ	TYR A		31.792	35.887	44.790	1.00 33.00	AAAA
ATOM	131	OH	TYR A		32.115	34.814	45.584	1.00 33.29	
MOTA	132	С	TYR A	17	31.296	41.000	40.613	1.00 44.43	AAAA

#### 218/263 Figure 19-3

MOTA	13	3 0	TYI	R A	17	31.346	42.194	40.905	1.0	0 44.68	AAAA
MOTA	134	1 N	PRO	) A	18	30.799	40.574	39.440		0 45.95	
MOTA	13	5 CI	PRO	) A	18	30.707	39.175	38.994		0 46.08	
MOTA	13	5 ÇA	PRO	) A	18	30.268	41.465	38.402		0 47.24	AAAA
MOTA	131	7 CE	PRO	) A	18	29.854	40.482	37.312	1.0	0 47.69	AAAA
MOTA	138	G CG	PRC	) A	18	30.876	39.338	37.511		0 46.79	AAAA
ATOM	139	) C	PRC	) A	18	29.129				0 48.98	
MOTA	140	0	PRO	) A	18	28.298				0 49.11	. AAAA
MOTA	141	N	LYS	A	19	29.114	43.593			0 50.59	AAAA
MOTA	142	CA	LYS	A	19	28.125				52.10	AAAA
ATOM	143	CB	LYS	Α	19	27.876				54.41	AAAA
ATOM	144	CG	LYS	Α	19	29.120				57.78	AAAA
MOTA	145	CD	LYS	Α	19	28.747				59.34	AAAA
ATOM	146	CE	LYS	A	19	29.978				60.33	AAAA
MOTA	147	NZ	LYS	Α	19	29.616				61.03	AAAA
ATOM	148	С	LYS	Α	19	26.764				51.53	AAAA
ATOM	- 149	.0	LYS	A	19	26.281				51.54	AAAA
ATOM	150	N	ASN	Α	20	26.146				50.13	AAAA
ATOM	151	CA	ASN	Α	20	24.831				48.44	AAAA
MOTA	152	CB	ASN	Α	20	24.336	42.061	37.209		49.67	AAAA
ATOM	153	CG.	ASN	A	20	25.389		36.613		51.61	AAAA
MOTA	154	OD:	1 ASN	Α	20	25.677	40.064	37.154		51.70	AAAA
ATOM	155	ND:	2 ASN	A	20	25.998	41.562	35.509		53.00	AAAA
ATOM	156	С	ASN	A	20	24.789	41.765	39.649	1.00	45.57	AAAA
ATOM	157	0	ASN	Α	20	23.764	41.127	39.877	1.00	44.67	AAAA
ATOM	158	N	HIS	A	21	25.883	41.662	40.398	1.00	42.71	AAAA
MOTA	159	CA	HIS		21	25.958	40.709	41.506	1.00	40.69	AAAA
MOTA	160	CB	HIS		21	27.216	39.857	41.353	1.00	40.16	AAAA
ATOM	161	CG	HIS		21	27.186	38.587	42.140	1.00	39.93	AAAA
MOTA	162		HIS		21	27.329	38.353	43.467	1.00	39.27	AAAA
MOTA	163		HIS		21	26.951	37.359	41.557		39.47	AAAA
ATOM	164		HIS		21	26.948	36.425	42.493		39.36	AAAA
ATOM	165		HIS		21	27.174	37.003	43.660		39.44	AAAA
MOTA	166	C	HIS		21	25.974	41.349	42.892		38.93	AAAA
ATOM	167	0	HIS		21	26.660	.42.338	43.116		38.78	AAAA
ATOM	168	N	PRO		22	25.229	40.778	43.853		37.11	AAAA
MOTA	169	CD	PRO		22	24.371	39.579	43.814		36.09	AAAA
MOTA	170 171	CA CB	PRO		22	25.224	41.361	45.199		35.81	AAAA
MOTA	172	CG	PRO PRO		22 22	24.473	40.306	46.012		36.04	AAAA
ATOM ATOM	173	C	PRO		22	23.464 26.638	39.810 41.637	45.003		36.19	AAAA
ATOM	174	Ö	PRO		22	26.867	42.653	45.751 46.417		34.39	AAAA
ATOM	175	N	LEU		23	27.572	40.731	45.451		34.09 31.98	AAAA
ATOM	176	CA	LEU		23	28.954	40.731	45.900		29.65	AAAA
ATOM	177	СВ	LEU		23	29.564	39.432	46.014		27.88	AAAA
ATOM	178	CG	LEU		23	28.896	38.528	47.048		27.31	AAAA AAAA
ATOM	179		LEU		23	29.656	37.217	47.149		26.64	. AAAA
ATOM	180		LEU		23	28.879	39.212	48.399		26.75	AAAA
ATOM	181	C	LEU		23	29.838		45.018		29.20	AAAA
ATOM	182	0	LEU .		23	31.057	41.606	45.028		28.38	AAAA
MOTA	183	N	LYS .		24	29.204	42.582	44.259		29.27	AAAA
ATOM	184	CA	LYS .		24	29.903	43.512			29.30	AAAA
ATOM	185	CB	LYS .	A.	24	28.881	44.091	42.405		29.75	AAAA
ATOM	186	CG	LYS A	A.	24	29.328	45.265	41.601		32.55	AAAA
ATOM	187	CD	LYS 2		24	28.537	46.526	41.994		34.37	AAAA
MOTA	188	CE	LYS A		24	27.025	46.337	41.835		34.32	AAAA
ATOM	189	NZ	LYS A		24	26.221	47.542	42.208		34:37	AAAA
ATOM	190	C	LYS A		24	30.580	44.620	44.224		28.14	AAAA
ATOM	191	0	LYS A		24	31.617	45.162	43.840		27.93	AAAA
ATOM	192	N	ILE A		25	29.990	44.919	45.377		27.07	AAAA
ATOM	193	CA	ILE A		25	30.468	45.945	46.296		25.82	AAAA
ATOM	194	CB	ILE ?		25	29.425	46.262	47.364		25.37	AAAA
ATOM	195		ILE A		25	28.190	46.846	46.737		25.71	AAAA
MOTA	196		ILE A		25	29.142	44.979	48.157		25.26	AAAA
MOTA	197	CD1	ILE A	. :	25	28.318	45.163	49.413	1.00	25.17	AAAA
MOTA	198	С	ILE A	. :	25	31.700	45.550	47.095		25.28	. AAAA

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- 51016	100	^	ILE A	25	32.037	44.379	47.183	1.00 24.48	AAAA ·
MOTA	199	0					47.714	1.00 24.98	AAAA
MOTA	200	И	PRO A	26	32.375	46.547			
ATOM	201	CD	PRO A	26	32.062	47.980	47.638	1.00 24.98	AAAA
ATOM	202	CA	PRO A	26	33.570	46.367	48.543	1.00 24.44	AAAA
		CB	PRO A	26	34.094	47.792	48.701	1.00 24.75	AAAA
MOTA	203							1.00 25.51	AAAA
ATOM	204	CG	PRO A	26	33.435	48.546	47.538		
MOTA	205	Ç	PRO A	26	33.021	45.838	49.862	1.00 23.42	AAAA
ATOM	206	0	PRO A	26	31.930	46.233	50.272	1.00 22.12	AAAA
					33.754	44.960	50.532	1.00 23.06	AAAA
MOTA	207	N	ARG A	27					
ATOM	208	CA	ARG A	27	33.244	44.421	51.776	1.00 23.04	AAAA
ATOM	209	CB	ARG A	27	32.633	43.043	51.492	1.00 22.20	AAAA
	210	CG	ARG A	27	31.463	43.152	50.503	1.00 19.84	AAAA
ATOM					30.762	41.844	50.160	1.00 18.64	AAAA
ATOM	211	CD	ARG A	27					
MOTA	212	NE	ARG A	27	30.181	41.168	51.315	1.00 16.51	AAAA
ATOM	213	CZ	ARG A	27	30.774	40.188	51.982	1.00 16.57	AAAA
	214		ARG A	27	31.969	39.763	51.605	1.00 17.50	AAAA '
ATOM				27	30.185	39.643	53.038	1.00 16.45	AAAA
ATOM	215		ARG A						
MOTA	216	C	ARG A	27	34.265	44.381	52.905	1.00 23.62	AAAA
MOTA	217	0	ARG A	27	34.107	45.077	53.919	1.00 23.69	AAAA
		N	VAL A	28	35.305	43.570	52.736	1.00 24.25	AAAA
MOTA	218					43.466	53.737	1.00 23.36	AAAA
ATOM	219	CA	VAL A	28	36.355				
ATOM	220	CB	VAL A	28	37.022	42.062	53.671	1.00 22.75	AAAA
ATOM	221	CG1	VAL A	28	38.292	42.031	54.475	1.00 22.95	AAAA
	222		VAL A	28	36.061	41.011	54.249	1.00 22.20	AAAA
MOTA					37.363	44.609	53.511	1.00 23.70	AAAA
ATOM	. 223	С	VAL A	28					
MOTA	224	0	VAL A	28	37.943	45.156	54.455	1.00 22.62	AAAA
MOTA	225	N	SER A	29	37.538	44.989	52.253	1.00 24.27	AAAA
	226	CA	SER A	29	38.444	46.082	51.910	1.00 26.03	AAAA
MOTA					38.632	46.178	50.381	1.00 25.95	AAAA
MOTA	227	CB	SER A	29					
ATOM	228	OG	SER A	29	37.395	46.417	49.716	1.00 27.57	AAAA
MOTA	229	C	SER A	29	37.793	47.354	52.440	1.00 25.52	aaaa
	230	Ó	SER A	29	38.463	48.311	52.828	1.00 25.49	AAAA
ATOM			LEU A	30	36.468	47.342	52.448	1.00 26.09	AAAA
MOTA	231	N					52.926	1.00 26.39	AAAA
MOTA	232	CA	LEU A	30	35.692	48.471			
MOTA	233	CB	LEU A	30	34.262	48.365	52.393	1.00 25.89	AAAA
ATOM	234	CG	LEU A	30	33.265	49.470	52.755	1.00 27.15	AAAA
			LEU A	30	32.486	49.101	53.999	1.00 26.34	AAAA
MOTA	235					50.813	52.897	1.00 25.81	AAAA
ATOM	236		LEU A	30	34.015				
MOTA	237	С	LEU A	30	35.713	48.534	54.453	1.00 26.26	AAAA
ATOM	238	0	LEU A	30	35.731	49.612	55.037	1.00 27.50	AAAA
	239	N	LEU A	31	35.730	47.379	55.097	1.00 25.57	AAAA
ATOM				31	35.776	47.343	56.545	1.00 26.87	AAAA
MOTA	240	CA	LEU A				57.029	1.00 27.28	AAAA
MOTA	241	CB	LEU A	31	35.752	45.900			
MOTA	242	CG	LEU A	31	35.135	45.563	58.383	1.00 27.87	AAAA
ATOM	243	CD1	LEU A	31	35.855	44.313	58.906	1.00 27.01	AAAA
			LEU A	31	. 35 261	46.706	59.372	1.00 26.32	AAAA
ATOM	244				37.087	48.003	57.012	1.00 28.08	AAAA
ATOM	245	С	LEU A	31					
ATOM	246	0	LEU A	31	37 094	48.854	57.901	1.00 27.42	AAAA
ATOM	247	N	LEU A	32	38.197	47.584	56.409	1.00 29.52	AAAA
	248	CA	LEU A	32	39.508	48.121	56.750	1.00 30.96	AAAA
MOTA					40.607	47.394	55.950	1.00 31.58	AAAA
ATOM	249	CB	LEU A	32			56.293	1.00 31.63	AAAA
ATOM	250	CG	LEU A	32	40.792	45.904			
ATOM	251	CD1	LEU A	32	41.810	45.246	55.380	1.00 31.31	AAAA
	252		LEU A	32	41.232	45.780	57.743	1.00 32.23	AAAA
ATOM							56.543	1.00 31.59	AAAA
ATOM	253	С	LEU A	32	39.599	49.635			
ATOM	254	0	LEU A	32	40.081	50.345	57.416	1.00 31.70	AAAA
	255	N	ARG A	33	39.140	50.129	55.398	1.00 32.72	AAAA
ATOM	•		ARG A	33	39.178	51.564	55.141	1.00 33.91	AAAA
ATOM	256	CA				51.903	53.743	1.00 35.10	AAAA
ATOM	257	CB	ARG A	33	38.643				
MOTA	258	CG	ARG A	33	39.627	51.609	52.621	1.00 37.84	AAAA
	259	CD	ARG A	33	39.310	52.412	51.374	1.00 39.33	AAAA
ATCM			ARG A	33	38.255	51.806	50.580	1.00 42.51	AAAA
MOTA	260	NE					49.541	1.00 44.22	AAAA
ATOM	261	CZ	ARG A	33	37.662	52.395	49.163		
ATOM	262	NHI	ARG A	33	38.016	53.617		1.00 44.61	AAAA
	263	NH2	ARG A	33	36.723	51.752	48.861	1.00 45.23	AAAA
ATOM			ARG A	33	38.352	52.305	56.168	1.00 33.48	AAAA
ATCM	264	С	א מיזע	73	عدد. در	JE.JUJ			•
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ATOM	265	0	ARG A	33		38.713	53.390	56.592	1.00 33.61	AAAA
MOTA	266	· N	PHE A	34		37.247	51.682	56.562	1.00 33.78	AAAA
	267		-	34		36.292	52.233	57.517		
MOTA		CA	PHE A						1.00 33.79	AAAA
MOTA	268	CB	PHE A	34		35.065	51.310	57.573	1.00 33.88	AAAA
MOTA	269	CG	PHE A	34		33.925	51.840	58.405	1.00 33.16	AAAA
MOTA	270		PHE A	34		33.108	52.856	57.925	1.00 32.77	AAAA
ATOM	271		PHE A	34		33.668	51.315	59.672	1.00 33.05	AAAA
MOTA	272	CE1	PHE A	34	•	32.044	53.343	58.695	1.00 32.86	AAAA
ATOM	273	CE2	PHE A	34		32.607	51.797	60.454	1.00 33.07	AAAA
		CZ	PHE A	34		31.794	52.809	59.966		
MOTA	274								1.00 32.58	AAAA
MOTA	275	С	PHE A	34	-	36.881	52.414	58.918	1.00 34.01	. AAAA
MOTA	276	0	PHE A	34		36.903	53.524	59.455	1.00 33.49	AAAA
	277	N	LYS A	35		37.350	51.324	59.516	1.00 34.00	AAAA
· ATOM										
MOTA	278	CA	LYS A	35		37.928	51.401	60.843	1.00 33.90	AAAA
ATOM	279	CB	LYS A	35		38.230	50.010	61.362	1.00 34.07	AAAA
ATOM	280	CG	LYS A	35		37.000	49.190	61.662	1.00 33.94	AAAA
-										
ATOM	281	CD	LYS A	35		37.414	47.810	62.106	1.00 35.31	AAAA `
ATOM	282	CE	LYS A	35		. 38.062	47.072	60.948	1.00 35.91	AAAA
MOTA	283	NZ	LYS A	35		39.058	47.928	60.236	1.00 36.19	AAAA
	284	C	LYS A	35		39.185	52.255	60.881	1.00 34.19	AAAA
ATOM										
MOTA	285	0 :	LYS A	35		39.554	52.775	61.929	1.00 34.32	AAAA
ATOM	286	N	ASP A	36		39.853	52.384	59.745	1.00 33.99	AAAA
ATOM	287	CA	ASP A	36		41.034	53.216	59.680	1.00 35.17	AAAA
	288		ASP A	36		41.812	52.943	58.388	1.00.37.40	AAAA
MOTA		CB								
ATOM	289	CG	ASP A	36		42.964	53.908	58.186	1.00 38.64	AAAA
ATOM	290	OD1	ASP A	36		43.648	54.218	59.184	1.00 40.02	AAAA
ATOM	291	002	ASP A	36		43.201	54.341	57.035	1.00 38.74	AAAA
			ASP A	36		40.568	54.670	59.724	1.00 35.85	AAAA
ATOM	292	С			•					
ATOM	293	0	ASP A	36		41.231	55.527	60.306	1.00 36.88	AAAA
ATOM-	294	N	ALA A	37		39.420€	54.940	59.111	1.00 34.96	AAAA
ATOM	295	CA	ALA A	37		38.851	56.280	59.108	1.00 34.47	AAAA
						37.751	56.373	58.067	1.00 33.80	
ATOM	296	CB	ALA A	37						AAAA
ATOM	297	С	ALAA	37		38.291	56.617	60.499	1.00 34.66	AAAA
ATOM	298	Q	ALA A	37		38.268	57.779	60.899	1.00 34.55	AAAA
ATOM	299	N	MET A	38		37.830	55.600	61.226	1.00 34.24	AAAA
						37.287	55.794	62.572	1.00 33.07	AAAA
MOTA	300	CA	MET A	38						
MOTA	301	CB	MET A	38		36.289	54.687	62.917	1.00 32.82	AAAA
ATOM	302	CG	MET A	38		35.084	54.559	61.996	1.00 32.72	AAAA
ATOM	303	SD	MET A	38		33.980	55.948	62.101	1.00 33.65	AAAA
				38		33.550	55.878	63.849	1.00 33.77	AAAA
ATOM	. 304	CE	MET A							
ATOM .	305	С	MET A	38		38.430	55.724	63.583	1.00 33.12	AAAA
ATOM	306	0	MET A	38		38.226	55.930	64.777	1.00 32.82	AAAA
ATOM	307	N	ASN A	39		39.628	55.428	63.090	1.00 32.64	AAAA
									1.00 32.38	AAAA
ATOM	308	CA	ASN A	39		40.805	55.266	63.935		
ATOM	309	CB	ASN A	39		41.200	56.600	64.589	1.00 32.93	AAAA
MOTA	310	CG	ASN A	39		41.393	57.736	63.571	1.00 34.40	AAAA
ATOM	311		ASN A	39	•	42.180	57.624	62.630	1.00 34.98	AAAA
		-							1.00 33.52	
ATOM	312		ASN A	39		40.677	58.838	63.772		AAAA
MOTA	313	С	asn a	39	•	40.483	54.212	65.009	1.00 31.69	AAAA
ATOM	314	0	ASN A	39		·40.565	54.490	66.205	1.00 31.12	AAAA
				40		40.095	53.010	64.570	1.00 31.76	AAAA
MOTA	315	N	LEU A							
MOTA	316	CA	LEU A	40		39.750	51.898	65.474	1.00 32.48	AAAA
ATOM	317	CB	LEU A	40		38.259	51.559	65.386	1.00 32.55	AAAA
ATOM	.318	CG	LEU A	40		37.231	52.581	65.879	1.00 32.84	AAAA
						35.837	52.089	65.554	1.00 33.79	AAAA
ATOM	319		LEU A	40						
ATOM	320	CD2	LEU A	40		37.372	52.798	67.376	1.00 32.45	AAAA
MOTA	321	С	LEU A	40		40.555	50.628	65.187	1.00 32.92	AAAA
	322		LEU A	40		40.196	49.530	65.618	1.00 31.64	AAAA
MOTA		0							1.00 34.12	
ATOM	323	N	ILE A	41		41.652	50.794	64.464		AAAA
ATOM	324	CA	ILE A	41		42.508	49.680	64.116	1.00 36.07	AAAA
ATOM	325	CB	ILE A	41		42.017	48.991	62.811	1.00 35.51	AAAA
						42.070	49.952	61.636	1.00 33.37	AAAA
ATOM	326		ILE A	41					1.00 35.97	
ATOM	327		ILE A	41		42.898	47.790	62.480	1.00 33.37	AAAA
ATOM	328	CD1	ILE A	41		42.854	46.701	63.500	1.00 37.19	<b>AAAA</b>
MOTA	329	c	ILE A	41		43.921	50.226	63.916	1.00 38.85	AAAA
								63.413	1.00 38.98	AAAA
MOTA	330	0	ILE A	41		44.106	51.346	00.413	1.00 50.50	FEET 1

	224									
MOTA	331		AS	P A	42	44.914	49.446	64.329	1.00 40.61	AAAA
ATOM	332	CA	AŞ	PA	42	46.309	49.843	64.181	1.00 42.57	AAAA
ATOM	333	CB	AS	P A	42	46.973	50.021	65.553	1.00 42.42	
ATOM	334			P A						AAAA
						46.316	51.110	66.381	1.00 42.27	AAAA
MOTA	335		1 AS			46.227	52.250	65.883	1.00 41.20	AAAA
ATOM.	336	OD	2 AS	P A	42	45.891	50.833	67.526	1.00 43.36	AAAA
ATOM	337	C	AS	P A	42	47.011	48.752	63.392	1.00 44.05	
	338									AAAA
ATOM				PΑ		46.525	47.620	63.333	1.00 44.88	`AAAA
ATOM	339	N	GL.	ΙV	. 43	48.147	49.090	62.789	1.00 45.10	AAAA
ATOM	340	CA	_GL	UΑ	43	48.905	48.141	61.980	1.00 46.11	AAAA
ATOM	341			UΑ		50.172				
							48.796	61.454	1.00 46.89	AAAA
MOTA	342			UΑ	-	49.924	50.057	60.668	1.00 49.30	AAAA
MOTA	343	CD	GL	UΑ	43	51.187	50.580	60.028	1.00 49.67	AAAA
ATOM	344	OE:	1 GL	U A	43	51.760	49.839	59.201	1.00 50.60	
ATOM	345		2 GLI			51.601	51.714	60.349		AAAA
									1.00 49.60	AAAA
MOTA	346	С		JA		49.290	46.859	62.701	1.00 46.27	- AAAA
MOTA	347	0	GLI	JA	43	49.214	45.773	62.131	1.00 46.00	AAAA
MOTA	348	N	LY!	S A	44	49.708	46.986	63.954	1.00 46.52	AAAA
ATOM	349	CA		SA		50.135	45.832	64.730		
									1.00 46.31	. AAAA
ATOM	350	CB		S A		50.762	46.306	66.048	1.00 48.16	AAAA
MOTA	351	CG	LYS	5 A	44	51.977	47.215	65.799	1.00 51.59	AAAA
MOTA	352	CD	LYS	5 A	44	52.641	47.734	67.071	1.00 52.87	AAAA
ATOM	353	CE	LYS		44	53.851	48.601			
								66.727	1.00 53.34	AAAA
MOTA	354	NZ	LYS		44	54.615	49.033	67.936	1.00 53.45	AAAA
ATOM	355	С	LYS	5 A	44	49.029	44.828	64.996	1.00 44.74	AAAA
ATOM	356	0	LYS	S A	44	49.296	43.735	65.480	1.00 45.35	AAAA
ATOM	357	N	GLU		45	47.793	45.190	64.659	1.00 42.49	
										AAAA
ATOM	358	CA	GLU		45	46.638	44.320	64.894	1.00 40.54	AAAA
MOTA	359	CB	GLU	ΙĄ	45	45.493	45.125	65.517	1.00 40.55	AAAA
ATOM	360	CG	GLU	J A	45	45.788	45.731	66.882	1.00 38.87	AAAA
ATOM	361	CD	GLU	Δ 1	45	44.663	46.618	67.360	1.00 37.57	
	362									AAAA
MOTA			. GLU		45	44.383	47.631	66.693	1.00 36.29	AAAA
ATOM	363	OE2	GLU	A	45	44.056	46.300	68.399	1.00 38.44	AAAA
ATOM	364	С	GLU	A	45	46.126	43.648	63.630	1.00 39.15	AAAA
ATOM	365	0	GLU	Α	45	45.301	42.737	63.681	1.00 39.29	AAAA
ATOM	366	N	LEU		46	46.619	44.115	62.497	1.00 37.62	
										AAAA
ATOM	367	CA	LEU		46	46.219	43.589	61.211	1.00 35.88	AAAA
MOTA	368	CB	LEU		46	46.125	44.750	60.229	1.00 36.09	AAAA
MOTA	369	CG	LEU	Α	46	45.608	44.550	58.817	1.00 36.50	AAAA
ATOM	370	CD1	LEU	Α	46	44.182	44.021	58.843	1.00 36.66	AAAA
MOTA	371		LEU		46	45.646	45.893	58.113		
									1.00 35.85	AAAA
ATOM	372	C	LEU		46	47.211	42.542	60.714	1.00 34.97	AAAA
MOTA	373	0	LEU	Α	46	48.424	42.670	60.900	1.00 35.72	AAAA
ATOM	374	N	ILE	A	47	46.680	41.484	60.118	1.00 33.25	AAAA
ATOM	375	CA	ILE		47	47.497	40.411	59.560	1.00 30.92	
	376	CB	ILE							AAAA
ATOM					47	47.144	39.024	60.167	1.00 31.22	AAAA
ATOM	377	CG2	ILE	А	47	48.093	37.97C	59.640	1.00 28.55	AAAA
ATOM	378	CG1	ILE	A	47	47.220	39.063	61.694	1.00 32.04	AAAA
ATOM	379	CDI	ILE	A	47	48.596	39.241	62.242	1.00 34.13	AAAA
	380	C	ILE		47					
MOTA						47.138	40.381	58.076	1.00 29.70	AAAA
MOTA	381	0	ILE		47	45.956		57.714	1.00 28.42	AAAA
MOTA	382	N	LYS	A	48	48.150	40.380	57.221	1.00 28.78	AAAA
ATOM	383	CA	LYS		48	47.920	40.349	55.784	1.00 28.42	AAAA
	384	CB	LYS				40.727	55.055		
MOTA					48	49.203			1.00 27.53	AAAA
MOTA	385	CG	LYS		48	49.116	40.695	53.556	1.00 28.97	AAAA
ATOM	386	CD	LYS	A	48	50.464	41.104	52.941	1.00 29.67	AAAA
ATOM	387	CE	LYS	A	48	50.493	40.893	51.432	1.00 29.41	AAAA
	388	NZ	LYS		48	49.409	41.645	50.764	1.00 29.68	
ATOM										AAAA
ATOM	389	C	LYS		48	47.449	38.950	55.375	1.00 27.81	AAAA
ATOM	390	0	LYS	A	48	48.024	37.938	55.787	1.00 27.96	AAAA
ATOM	391	N	SER		49	46.385	38.892	54.581	1.00 26.82	AAAA
	392	CA	SER		49	45.854	37.611	54.141	1.00 26.41	
ATOM										AAAA
ATOM	393	CB	SER		49	44.514	37.795	53.420	1.00 25.40	AAAA
ATOM	394	OG	SER	A .	. 49	43.541	38.349	54.276	1.00 25.58	AAAA
ATOM	395	С	SER	A	49	46.814	36.891	53.207	1.00 26.03	AAAA
	396	ō	SER		49	47.462	37.513	52.373	1.00 26.98	
ATOM	330	J	المندن	A	37	47.402	دید. ار	26.313	1.00 20.38	AAAA
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				1 Iguic 17	,			
ATOM	397	N ARG A	¥ 50	46.910	35.576	53.354	1.00 25.51	аааа
ATOM		CA ARG A	A 50	47.755	34.794	52.474	1.00 25.45	AAAA
ATOM		CB ARG A		48.807	33.985	53.252	1.00 25.85	AAAA
ATOM		CG ARG A		48.229	32.819	54.009	1.00 27.16	AAAA
ATOM	•	CD ARG A		49.280	31.995		1.00 27.57	AAAA
MOTA MOTA		NE ARG A CZ ARG A		48.673 48.106	30.896 29.820	55.482 54.946	1.00 27.90 1.00 28.34	AAAA
ATOM		NH1 ARG A		48.055	29.672	53.630	1.00 28.19	AAAA AAAA
ATOM		NH2 ARG A		47.592	28.884	55.735	1.00 28.62	AAAA
ATOM		C ARG A		46.806	33.834	51.762	1.00 24.91	AAAA
ATOM	407	O ARG A	50	45.740	33.510	52.283		AAAA
MOTA		N PRO A		47.172	33.392	50.549	1.00 24.28	AAAA
ATOM		CD PRO A		48.361	33.761	49.770	1.00 24.13	AAAA.
MOTA ATOM		CA PRO A CB PRO A		46.355 47.012	32.462 32.512	49.776 48.390	1.00 24.18	AAAA
ATOM		CG PRO A		47.766	33.862	48.405	1.00 24.24	AAAA AAAA
MOTA		PRO A		46.473	31.070	50.393	1.00 23.69	AAAA
ATOM		PRO A		47.545	30.680	50.839	1.00 24.13	AAAA
ATOM	415 N	N ALA N	. 52	45.381	30.325	50.422	1.00 23.36	AAAA
ATOM		CA ALA A		45.419	28.972	50.952	1.00 23.64	AAAA
MOTA		B ALA A		44.012	28.405		1.00 23.86	AAAA
MOTA	418 C	ALA A ALA A		. 46.260 46.240	28.145	49.994	1.00 23.58 1.00 24.52	AAAA
ATOM ATOM	419 K	* Committee of the comm		47.009	28.383 27.185	48.806 50.501	1.00 24.52	AAAA A <b>AA</b> A
ATOM		A THR A		47.815	26.352	49.628	1.00 26.26	AAAA
MOTA		B THR A		48.933	25.642	50.405	1.00 26.37	AAAA
MOTA		G1 THR A		48.355	24.763	51.375	1.00 26.51	AAAA
MOTA		G2 THR A		49.810	26.648	51.106	1.00 24.48	AAAA
MOTA	425 C			46.889	25.299	49.034	1.00 27.63	AAAA
ATOM ATOM	426 O 427 N		53 54	45.870 47.240	24.982	49.620 47.867	1.00 29.22	AAAA AAAA
ATOM		A LYS A	54	46.450	23.752	47.189	1.00 30.61	AAAA
ATOM		B LYS A	54	47.249	23.182	46.015	1.00 31.68	AAAA
ATOM		G LYS A	54 .		22:020	45.304	1.00 34.38	AAAA
ATOM		D LYS A	54	45.449	22.464	44.417	1.00 36.00	AAAA
ATOM	432 C 433 N	E LYS A Z LYS A	54 54	45.943	22.850 21.664	43.025 42.236	1.00 37.55	AAAA
ATOM ATOM	433 N		54	46.425 46.127	22.640	48.170	1.00 37.57 1.00 31.26	AAAA AAAA
ATOM	435 0		54	45.025	22.097	48.176	1.00 31.72	AAAA
ATOM	436 N	GLU A	55	47.102	22.312	49.006	1.00 31.88	· AAAA
ATOM	437 C		55	46.961	21.260	50.011	1.00 32.29	AAAA
ATOM	438 C		55	48.266	21.089	50.778	1.00 34.43	AAAA
MOTA	439 C		55 55	48.265 49.513	19.901 19.839	51.706 52.584	1.00 38.39 1.00 41.46	AAAA
MOTA MOTA		E1 GLU A	55	49.745	18.770	53.200	1.00 41.40	AAAA AAAA ·
ATOM		E2 GLU A	55	50.245	20.859	52.672	1.00 42.45	AAAA
ATOM	443 C		55	45.851	21.555	51.013	1.00 30.43	AAAA
ATOM	444 0		55	45.048	20.681	51.332	1.00 30.59	AAAA
ATOM	445 N		56	45.822	22.782	51.517	1.00 28.23	AAAA
ATOM	446 CZ 447 CI		56 56	44.812 45.078	23.164	52.488 52.989	1.00 27.69 1.00 27.90	AAAA AAAA
MOTA MOTA	448 CC		56	46.434	24.721	53.670	1.00 27.90	AAAA
ATOM	449 CI		56	46.769	26.135	54.098	1.00 26.35	AAAA
ATOM		E1 GLU A	56	46.615	27.057	53.265	1.00 25.12	AAAA
ATOM	451 OF	E2 GLU A	56	47.213	26.315	55.255	1.00 25.70	AAAA
ATOM	452 C	GLU A	56	43.408	23.043	51.914	1.00 26.99	AAAA
ATOM	453 O	GLU A	56 57	42.495	22.574	52.588	1.00 26.25	AAAA
ATOM	454 N 455 CA	LEU A	57 57	43.252 41.965	23.447 23.389	50.659 49.967	1.00 27.26 1.00 27.17	AAAA AAAA
ATOM ATOM	456 CE		57	42.077	24.063	48.596	1.00 26.62	AAAA
ATOM	457 CG		57	42.491	25.545	48.656	1.00 27.64	AAAA
ATOM		1 LEU A	57	42.770	26.108	47.269	1.00 26.66	AAAA
MOTA	459 CD	2 LEU A	57	41.389	26.341	49.349	1.00 26.92	AAAA
ATOM	460 C	LEU A	57 52	41.552	21.946	49.796	1.00 27.26	AAAA
ATOM	461 O	LEU A	57 50	40.363	21.612	49.816	1.00 27.53	AAAA
MOTA	462 N	LEU A	58	42.547	21.085	49.641	1.00 27.42	<b>AAAA</b>
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#### Figure 19-8

ATOM	463	CA	LEU	Α	58		42.293	19.675	49.457	1.00	26.10	AAAA
ATOM	464	CB	LEU	A	58	4,	43.486	19.019	48.794		25.43	AAAA
MOTA	465		LEU		58		43.623	19.577	47.385		26.66	AAAA
ATOM	466		LEU		58		44.760	18.884	46.705		27.12	AAAA
ATOM	467	CD2	LEU	Α	58		42.334	19.355	46.600	1.00	26.43	AAAA
ATOM	468	С	LEU	Α	58		41.938	18.956	50.731		25.79	AAAA
ATOM	469	0	LEU	Α	58		41.648	17.763	50.692		26.50	AAAA
ATOM	470	N	LEU	A	59		41.977	19.666	51.858		24.91	AAAA
ATOM	471	CA	LEU	Α	59		41.595	19.070	53.136		25.15	AAAA
ATOM	472	CB	LEU		59		41.958	19.991	54.322		25.44	AAAA
ATOM	473	CG	LEU	A	59		43.423	20.280	54.710		24.67	AAAA
ATOM	474	CD1	LEU	A	59		43.502	21.461	55.652		23.70	AAAA
MOTA	475	CD2	LEU	Α	59		44.044	19.044	55.357		24.08	AAAA
ATOM	476	С	LEU	Α	59		40.074	18.870	53.090		25.41	AAAA
ATOM	477	0	LEU	Α	59		39.503	18.266	53.993		25.88	AAAA
ATOM	478	N	PHE	A	60		39.436	19.392	52.031		25.05	AAAA
ATOM	479	CA	PHE	Α	60		37.983	19.276	51.823		24.11	AAAA
ATOM	480	CB	PHE	A	60		37.250	20.476	52.440		21.80	. AAAA
ATOM	481	CG	PHE	A	60		35.778	20.534	52.098		20.07	AAAA
MOTA	482	CD1	PHE	Α	60		34.917	19.501	52.462		19.27	AAAA
MOTA	483	-CD2	PHE	Α	60		35.249	21.628	51.399		19.82	AAAA
ATOM	484	CE1	PHE	Α	60		33.550	19.557	52.136	1.00	19.26	AAAA
MOTA	485	CE2	PHE	Α	60		33.890	21.688	51.071		17.45	AAAA
ATOM	486	CZ	PHE	Α	60		33.042	20.652	51.440	1.00	17.92	AAAA
MOTA	487	С	PHE	Α	60		37.557	19.139	50.345	1.00	24.02	AAAA
MOTA	488	0	PHE	A	60		36.846	18.201	49.974	1.00	23.27	AAAA
MOTA	489	N	HIS	Α	61		37.982	20.079	49.511	1.00	24.40	AAAA
MOTA	490	CA	HIS		61		37.626	20.053	48.099		25.04	AAAA
ATOM	491	CB	HIS		61		37.768	21.449	47.494	1.00	24.19	AAAA
MOTA	492	CG	HIS		61		36.744	22.429	47.979	1.00	24.44	AAAA
MOTA	493		HIS		61		35.429	22.559	47.683	1.00	24.12	AAAA
MOTA	494		HIS		61		37.038	23.444	48.864		24.36	AAAA
ATOM	495		HIS		61		35.952	24.159			23.18	AAAA
MOTA	496		HIS		61		34.962	23.643	48.385		23.91	AAAA
ATOM	497	C	HIS		61		38.416	19.054	47.253		25.60	AAAA
ATOM	498	0	HIS		61		39.596	18.805	47.498		26.94	AAAA
ATOM	499	N	THR		62		37.754	18.496	46.244		26.68	AAAA
ATOM	500 501	CA CB	THR		62		38.369 37.290	17.522	45.333		28.17	AAAA
ATOM ATOM	502		THR THR		62 62 .		36.544	16.695 17.541	44.614 43.731		28.15	AAAA
ATOM	503		THR		62		36.334	16.094	45.629		28.24	AAAA
ATOM	504	C	THR		62		39.226	18.217	44.278		29.28	AAAA AAAA
ATOM	505	ŏ	THR		62		38.876	19.286	43.792		29.52	AAAA
ATOM	506	N	GLU		63		40.344	17.606	43.912		31.33	AAAA
ATOM	507		GLU		63		41.249	18.202	42.928		32.42	AAAA
ATOM	508		GLU		63		42.333	17.219	42.536			AAAA
ATOM	509		GLU		63		43.304	16.869	43.609	1 00	37.20	AAAA
ATOM	510		GLU .		63		44.427	16.022	43.052		38.79	AAAA
ATOM	511		GLU .		63		45,100	16.499	42.097		37.96	AAAA
ATOM	512	OE2	GLU :	A	63		44.619	14.892	43.564		39.68	AAAA
MOTA	513	С	GLU :	A	63		40.607	18.687	41.639		31.96	AAAA
ATOM	514	0	GLU .	A	63		40.824	19.816	41.215	.1.00	32.10	AAAA
ATOM	515		ASP A		64		39.845	17.814	40.998		31.52	AAAA
ATOM	516	CA .	ASP A	A	64		39:204	18.165	39.753	1.00	31.36	AAAA
ATOM	517	CB .	ASP A	A	64		38.301	17.018	39.295	1.00	33.99	AAAA
ATOM	518	CG .	ASP A	Ą	64		37.213	16.694	40.302	1.00	37.38	AAAA
MOTA	519	OD1	ASP A	A.	64		36.375	15.801	40.027	1.00	39.80	AAAA
MOTA	520	OD2			64		37.188	17.332	41.374	1.00	38.67	AAAA
MOTA	521	C Z	ASP I	Ą	64		38.412	19.465	39.902		30.02	AAAA
ATOM	522		ASP A		64		38.462	20.331	39.026	1.00	30.47	AAAA
ATOM	523	N :	ryr 2	4	65		37.695	19.608	41.012	1.00	27.51	AAAA
ATCM	524		ryr 1		65		36.918	20.814	41.248		26.03	AAAA
ATOM	525		ryr 1		65		36.010	20.654	42.467		25.42	AAAA
MOTA	526.		ryr 1		65		35.339	21.946	42.866		24.90	AAAA
МОТА	527	CD1			65		34.525	22.636	41.964		25.04	AAAA
ATOM	528	CE1	ryr A	1	65		33.914	23.823	42.308	1.00	25.01	AAAA.

MOTA	529	CD	2 TYR	A 65		35.525	22.486	44.136	1.00	24.65		AAAA
MOTA	530	) CE	2 TYR	A 65		34.920	23.677	44.497	1.00	25.86		AAAA
MOTA	531	. CZ	TYR	A 65		34.110	24.349	43.576				AAAA
MOTA	532	OH	TYR	A 65		33.499	25.543	43.924	1.00	27.20		AAAA
MOTA	533	C	TYR	A 65		37.814			1.00 2			AAAA
MOTA	534	0	TYR	A 65		37.460	23.129	41.096	1.00 2	25.62		AAAA
MOTA	535	N	ILE	A 66		38.965			1.00 2			AAAA
ATOM	536	CA	ILE	A 66	•	39.877	22.902	42.328	1.00			AAAA
MOTA	537	CB	ILE	A 66		40.924	22.520	43.402	1.00			AAAA
ATOM	538	CG	2 ILE	A 66		41.927	23.652		1.00 2			AAAA
MOTA	539	CG	1 ILE	A 66		40.220			1.00 2			AAAA
MOTA	540	CD	1 ILE	A 66		39.528			1.00		•	AAAA.
MOTA	541	С	ILE .	A 66		40.558			1.00 2			AAAA
MOTA	542	0	ILE .	A 66		40.636	24.425	40.665	1.00 2			AAAA
MOTA	543	N	ASN .	A 67		41.036	22.262	40.295	1.00 2			AAAA
ATOM	544	CA	ASN .	A 67		41.698	22.545	39.02 <del>9</del>	1.00 2	3.92		AAAA
ATOM	545	CB	ASN .	A 67		42.292	21.261	38.395	1.00 2			AAAA
MOTA	546	CG	ASN A	A 67		43.344	20.588	39.289	1.00 2			AAAA
MOTA	547	OD:	L ASN	A 67		44.196	21.256	39.859	1.00 2	3.47		AAAA
ATOM	548	ND2	ASN A	A 67.		43.290	19.258	39.392	1.00 2	3.20		AAAA
MOTA	549	C.	ASN A	A 67		40.717	23.216	38.063	1.00 2	3.82		AAAA
MOTA	550	0	ASN A	A '67		41.123	23.996	37.204	1.00 2	4.63		AAAA
ATOM	5 <b>5</b> 1	N	THR A	A 68		39.427	22.928	38.213	1.00 2	4.08		AAAA
ATOM	552	CA	THR A			38.428	23.534	37.343	1.00 2	5.28		AAAA
MOTA	553	CB	THR A	4 68		37.030	22.904	37.525	1.00 2	4.55		AAAA
ATOM	554	OG1				37.090	21.500	37.258	1.00 2	4.64		AAAA
ATOM	555	CG2	THR 2	4 68		36.049	23.534	36.564	1.00 2	3.58		AAAA
MOTA	556	С	THR. A	A 68		38.322	25.023	37.664	1.00 2	6.31		AAAA
MOTA	557	0	THR ?			38.114	25.854	36.771	1.00 2	6.69		AAAA
ATOM	558	Ŋ	LEU A			38.462	25.351	38.945	1.00 2			AAAA
ATOM	559	CA	LEU A			38.381	26.729	39.378	1.00 2			AAAA
ATOM	560	CB	LEU A			38.321	26.807	40.904	1.00 2			AAAA
MOTA	561	CG	LEU A			37.003	26.397	41.551	1.00 2			AAAA
MOTA	562		LEU A			37.088	26.491	43.062	1.00 2			AAAA
ATOM	563		LEU A			35.933	27.316	41.044	1.00 2			AAAA
ATOM	564	C	LEU A			39.570	27.508	38.867	1.00 2			AAAA
MOTA	565	0	LEU A			39.425	28.619	38.356	1.00 2			AAAA
ATOM	566 567	N CA	MET A			40.748	26.914	39.009	1.00 2			AAAA
ATOM	568	CB	MET A			41.981	27.536 26.692	38.571 39.044	1.00 2			AAAA
ATOM ATOM	569	CG	MET A			43.164	26.528	40.562	1.00 3			AAAA
ATOM	570	SD	MET A			44.608	25.684	41.183	1.00 3			AAAA
ATOM	571	CE	MET A			45.859	26.820	40.670	1.00 3			AAAA
ATOM	572	C	MET A			42.017	27.723	37.057	1.00 3			AAAA AAAA
ATOM	573	ō	MET A			42.462	28.769	36.559	1.00 3			AAAA
MOTA		N	GLU A			41.538	26.719	36.328	1.00 3			AAAA
ATOM	575	CA	GLU A			41.519	26.795	34.874	1.00 3			AAAA
ATOM	576	CB	GLU A			41.140	25.442	34.266	1.00 3			AAAA
ATOM	577	CG	GLU A			41.122	25.430	32.731	1.00 3			AAAA
ATOM	578	CD	GLU A	71		42.513	25.676	32.093	1.00 4			AAAA
ATOM	579	OE1	GLU A	71		42.570	25.798	30.844	1.00 4			AAAA
ATOM	580	OE2	GLU A	71		43.541	25.738	32.825	1.00 40			AAAA
ATOM	581	C	GLU A	71		40.537	27.851	34.392	1.00 29			AAAA
MOTA	582	0	GLU A	71		40.852	28.642	33.508	1.00 27			AAAA
ATOM	583	N	ALA A	72		39.352	27.855	34.992	1.00 29			AAAA
ATOM	584	CA	ALA A	72	:	38.296	28.790	34.635	1.00 29			AAAA
ATOM	585	CB	ALA A	72	:	37.022	28.432	35.374	1.00 29			AAAA
ATOM	586	С	ALA A	72.	:	38.667	30.238	34.907	1.00 30			AAAA
MOTA	587	0	ALA A	72	3	38.359	31.122	34.108	1.00 31	.27		AAAA
ATOM	588	N	GLU A	73	-	39.336	30.491	36.023	1.00 31			AAAA
MOTA	589	CA	GLU A	73		9.710	31.856	36.346	1.00 31	. 65		AAAA
MOTA	590	CB	GLU A	73	4	10.243	31.954	37.785	1.00 30	.52		AAAA
MOTA	591		GLU A	73		10.643	33.370	38.198	1.00 28			AAAA
MOTA	592		GLU A	73		1.076	33.484	39.651	1.00 28			AAAA
MOTA	593		GLU A	73		10.239	33.260	40.546	1.00 28	.94		AAAA
ATOM	594	OE2	GLU A	73	. 4	2.258	33.795	39.906	1.00 28	.57		AAAA

ATOM	595	С	GLU A	73	40.726	32.461	35.378	1.00 33.54	AAAA
ATOM	596	Q	GLU A	73	40.456	33.499	34.767	1.00 34.93	AAAA
ATOM	597	N	ARG A	74	41.885	31.832	35.214	1.00 34.35	AAAA
ATOM	598	ĊA	ARG A	74	42.890	32.428	34.334	1.00 36.04	AAAA
ATOM	599	CB	ARG A	74	44.238	31.710	34.482	1.00 36.92	AAAA
ATOM-	600	CG	ARG A	74	44.327	30.313	33.923	1.00 38.14	AAAA
	601	CD	ARG A	74	45.508	29.589	34.543	1.00 39.55	AAAA
ATOM	602	NE	ARG A	74	45.893	28.404	33.785	1.00 42.02	AAAA
ATOM	603	CZ	ARG A		46.632	28.436	32.675	1.00 42.69	AAAA
ATOM			-ARG A	74	47.071	29.593	32.191	1.00 42.76	AAAA
ATOM	604			74	46.933	27.309	32.046	1.00 42.78	AAAA
MOTA	605		ARG A	74		32.532		1.00 42.92	AAAA
MOTA	606	C	ARG A	74	42.476		32.864 32.187		
ATOM	607	0	ARG A	74	42.842	33.493	32.167	1.00 37.73	AAAA
ATOM	608	N	SER A	75	41.711	31.567		1.00 36.60	AAAA
ATOM	609	CA	SER A	75	41.248	31.622	30.987	1.00 36.82	AAAA
MOTA	610	СВ	SER A	75	40.916	30.218	30.478	1.00 36.10	- AAAA
ATOM	611	OG	SER A	75	39.736	29.723	31.083	1.00 36.39	AAAA
MOTA	612	С	SER A	75	39.980	32.476	31.001	1.00 36.90	AAAA
ATOM	613	0	SER A	75	39.401	32.791	29.963	1.00 36.25	AAAA
ATOM	614	N	GLN A	76	39.568	32.845	32.208	1.00 37.62	AAAA
ATOM	615	CA	GLN A	76	38.368	33.639	32.427	1.00 37.92	AAAA
ATOM	616	CB	GLN A	76	38.613	35.100	32.049	1.00 38.23	AAAA
ATOM	617	CG	GLN A	76	37.630	36.048	32.717	1.00 40.67	, AAAA
ATOM	618	CD	GLN A	76	37.929	36.298	34.199	1.00 41.40	AAAA
ATOM	619	OE1	GLN A	76	38.226	35.379	34.973	1.00 40.79	AAAA
ATOM	620	NE2	GLN A	76	37.833	37.556	34.597	1.00 42.32	AAAA
ATOM	621	С	ĠŁN A	76	37.223	33.064	31.600	1.00 37.75	AAAA
ATOM	622	0	GLN A	76	- 36.521	33.789	30.901	1.00 38.13	AAAA
MOTA	623	N	SER A	.77	37.045	31.749	31.685	1.00 37.52	AAAA
ATOM	624	CA	SER A	77	35.990	31.061	30.950	1.00 37.75	AAAA
ATOM	625	CB	SER A	77	36.537	30.440	29.664	1.00 37.90	· AAAA
ATOM	626	OG	SER A	77	36.851	31.441	28.724	1.00 40.32	AAAA
ATOM	627	С	SER A	77	35.338	29.960	31.757	1.00 37.55	AAAA
MOTA	628	0	SER A	77	35.790	29.620	32.846	1.00 36.81	AAAA
ATOM	629	N	VAL A	78	34.264	29.412	31.198	1.00 37.82	AAAA
MOTA	630	CA	VAL A	78	33.538	28.309	31.812	1.00 37.99	AAAA
ATOM	631	CB	VAL A	78	32.027	28.514	31.715	1.00 37.19	AAAA
MOTA	632		VAL A	78	31.310	27.439	32.497	1.00 36.84	AAAA
ATOM	633		VAL A	78	31.662	29.906	32.201	1.00 37.60	AAAA
ATOM	634	С	VAL A	78	33.918	27.089	30.976	1.00 38.28	AAAA
ATOM	635	0	VAL A	78	33.497	26.959	29.819	1.00 39.18	AAAA
ATOM	636	N	PRO A	79	34.734	26.187	31.537	1.00 37.69	AAAA
ATOM	637	CD	PRO A	79	35.347	26.1 <b>67</b>	32.869	1.00 37.65	AAAA
ATOM	638	CA	PRO A	79	35.146	24.998	30.797	1.00 37.54	AAAA
ATOM	639	CB	PRO A	79	36.127	24.325	31.759	1.00 37.45	AAAA
TOM	640	CG	PRO A	79	36.655	25.489	32.557	1.00 37.65	AAAA
ATOM	641	C	PRO A	79	33.980	24.089	30.434	1.00 37.20	* AAAA
ATOM	642	0	PRO A	79	32.958	24.050	31.120	1.00 36.43	AAAA
ATOM	643	N	LYS A	80	34.154	23.363	29.338	1.00 37.42	AAAA
ATOM	644	CA	LYS A	80	33.160	22.423	28.855	1.00 37.35:	AAAA
ATOM	645	CB	LYS A	80	33.757	21.586	27.725	1.00 37.99	AAAA
ATOM	646	CG	LYS A	80	32.928	20.379	27.280	1.00 38.94	AAAA
ATOM	647	CD	LYS A	80	31.835	20.710	26.286	1.00 39.07	AAAA
ATOM	.648	CE	LYS A	80	31.320	19.402	25.688	1.00 40.43	AAAA
ATOM	649	NZ	LYS A	80	30.498	19.543	24.450	1.00 40.48	AAAA
ATOM	650	C	LYS A	80	32.752	21.515	30.003	1.00 36.85	AAAA
	651	ŏ	LYS A	80	33.610	20.942	30.676	1.00 36.56	AAAA
ATOM	652	N	GLY A	81	31.443	21.408	30.217	1.00 35.94	AAAA
ATOM	653	CA	GLY A	81	30.903	20.570	31.268	1.00 35.48	AAAA
ATOM		C	GLY A	81	31.110	21.054	32.693	1.00 35.23	AAAA
ATOM	654 655	0	GLY A	81	30.749	20.355	33.644	1.00 35.46	AAAA
FIOM	655 656		ALA A	82	31.677	22.241	32.867	1.00 35.17	AAAA
ATOM	656 657	N	ALA A	82	31.919	22.743	34.213	1.00 35.02	AAAA
ATOM	657	CA	ALA A	82	33.076	23.743	34.208	1.00 35.13	AAAA
ATOM	658	CB.	ALA A	82	30.674	23.378	34.797	1.00 34.39	AAAA
ATOM	659	C O	ALA A	82	30.451	23.378	36.001	1.00 33.82	AAAA
ATOM	660	•	ALIA II		20.431		•	_	•

### ; **226/263** Figure 19-11

ATOM	661	N	ARG	Α	83		29.858	23,960	33.932	1.00 34.77	AAAA
ATOM	662	ÇA	ARG	Α	83		28.637		34.361	1.00 35.34	AAAA
ATOM	663	CB	ARG		83		27.899		33.150	1.00 36.26	
ATOM	664	CG	ARG		83		27.045	-	33.464	1.00 37.09	AAAA
ATOM	665	CD	ARG		83		26,209		34.686		AAAA
	666									1.00 37.48	AAAA
ATOM		NE	ARG		83		25.475		35.134	1.00 37.35	AAAA
MOTA	667	CZ	ARG		83		24.711		36.218	1.00 37.77	. AAAA
ATOM	668		L ARG		83		24.606		36.940	1.00 37.29	AAAA
ATOM	669	NH2	2 ARG	A.	. 83		24.040	28.401	36.568	1.00 38.34	AAAA
MOTA	670	С	ARG	Α	83		27.739	23.603	35.065	1.00 36.30	AAAA
MOTA	671	0	ARG	Α	83		27.232	23.854	36.154	1.00 36.17	AAAA
MOTA	672	N	GLU	Α	84		27.565	22.450	34.431	1.00 37.19	AAAA
ATOM	673	CA	GLU	A	84		26.721	21.382	34.948	1.00 37.80	AAAA ·
ATOM	674	CB	GLU	A	84		26.466	20.375	33.833	1.00 40.55	AAAA
ATOM	675	CG	GLU	Α	84		25.643	19.171	34.232	1.00 43.12	AAAA
MOTA	676	CD	GLU	А	84		25.362	18.268	33.046	1.00 44.98	AAAA
ATOM	677		GLU		84		24.573	17.301	33.195	1.00 46.36	AAAA
ATOM	678		GLU		84		25.937	18.532	31.962	1.00 44.94	
`ATOM	679	C	GLU		84		27.290	20.657	36.158	1.00 37.07	AAAA
	680	ō	GLU		84		26.642	20.555	37.199		AAAA
ATOM	681	И	LYS		85					1.00 36.17	AAAA
MOTA							28.506	20.152	35.999	1.00 36.23	AAAA
ATOM	682	CA	LYS		85		29.202	19.412	37.043	1.00 35.36	AAAA
ATOM	683	CB	LYS		85		30.449	18.761	36.437	1.00 36.96	AAAA
MOTA	584	CG	LYS		85		31.394	18.158	37.465	1.00 39.04	AAAA
MOTA	685	CD	LYS		85	,	30.995	16.766	37.919	1.00 40.59	AAAA
MOTA	686	CE	LYS		85		31.508	15.719	36.933	1.00 41.88	AAAA
ATOM	687	NZ	LYS		85		32.998	15.817	36.757	1.00 42.00	AAAA
MOTA	688	С	LYS		85		29.620	20.202	38.289	1.00 33.86	AAAA
ATOM	689	0	LYS	A	85		29.576	19.679	39.404	1.00 33.82	AAAA
ATOM	690	N	TYR	Α	86		30.014	21.458	38.097	1.00 32.06	AAAA
MOTA	691	CA	TYR	A	86		30.514	22.279	39.194	1.00 29.44	AAAA
ATOM	692	CB	TYR	Α	86		31.956	22.683	38.875	1.00 29.97	AAAA
MOTA	693	CG	TYR	Α	86		32.872	21.496	38.621	1.00 29.99	AAAA
ATOM	694	CD1	TYR	A	86	•	33.281	20:666	39.666	1.00 29.24	AAAA
ATOM .	695	CE1	TYR	A	86		34.126	19.582	39.437	1.00 29.85	AAAA
ATOM	696	CD2	TYR	Α	86		.33.329	21.204	37.329	1.00 30.16	AAAA
ATOM	697	CE2	TYR	А	86		34.173	20.118	37.087	1.00 29.61	AAAA
ATOM	698	CZ	TYR	Α	86	•	34.570	19.313	38.148	1.00 29.79	AAAA
ATOM	699	ОН	TYR		86		35.414	18.253	37.923	1.00 29.48	AAAA
ATOM	700	C	TYR		86		29.705	23.509	39.572	1.00 27.81	AAAA
ATOM	701	ō	TYR		86		30.052	24.202	40.524	1.00 27.56	AAAA
ATOM	702	N	ASN .		87		28.642	23.784	38.828	1.00 26.60	AAAA
ATOM	703	CA	ASN		87		27.777	24.924	39.111	1.00 26.56	AAAA
ATOM	704	CB	ASN		87		27.172	24.772	40.508	1.00 26.39	
ATOM	705	CG	ASN .		87		25.863	25.544	40.684	1.00 26.64	AAAA
	706		ASN .		87			25.632	41.790	1.00 26.84	AAAA
ATOM	707				87		25.335				AAAA
ATOM			ASN				25.330	26.084	39.597	1.00 26.33	AAAA
ATOM	708	C	ASN .		87		28.587	26.217	39.024	1.00 26.40	AAAA
MOTA	709	0	ASN .		87		28.430	27.129	39.832	1.00 24.80	AAAA
ATOM	710	N	ILE A		88			. 26.273	38.015	1.00 27.57	AAAA
MOTA	711	CA	ILE A		88		30.330	27.409	37.767	1.00 27.88	AAAA
ATOM	712	CB	ILE A		88		31.817	26.932	37.648	1.00 27.38	AAAA
ATOM	713	CG2	ILE A		88		32.684	27.994	36.986	1.00 26.34	AAAA
MOTA	714	CG1	ILE 2	A.	88		32.354	26.543	39.026	1.00 28.35	AAAA
MOTA	715	CD1	ILE A	Ą	88		32.356	27.671	40.042	1.00 27.78	AAAA
ATOM	716	С	ILE A	A.	88		29.946	28.110	36.472	1.00 29.17	· AAAA
ATOM	717	0	ILE 2	A.	88		29.530	27.469	35.515	1.00 29.75	AAAA
ATOM	718	N	GLY A		89		30.092	29.429	36.443	1.00 29.96	AAAA
ATOM	719		GLY 2		89		29.791	30.162	35.229	1.00 30.24	AAAA
ATOM	720		GLY ?		89.		28.430	30.805	35.242	1.00 30.44	AAAA
ATOM	721		GLY A		89		28.177	31.769	34.514	1.00 31.14	AAAA
ATOM	722		GLY A		90		27.542	30.268	36.061	1.00 30.00	AAAA .
	723		GLY A		90		26.221	30.841	36.129	1.00 30.52	AAAA .
ATOM									36.661	1.00 30.52	
ATOM	724		GLY A		90		26.283	32.262			AAAA
MOTA	725		GLY A		90		27.356	32.795	36.962	1.00 30.34	AAAA
ATOM	726	N '	TYR A		91		25.112	32.873	36.768	1.00 31.09	AAAA



ATOM	727	CA	ጥህነ	R A	91	24.977	34.213	37.290	1.00 31.27	2222
	728	CB								AAAA
ATOM			-	R A		23.515	34.634	37.195	1.00 31.82	AAAA
ATOM	729	CG		R A		23.169	35.825	38.047	1.00 31.81	AAAA
ATOM	730	CD:	l TY	RA	91	23.536	37.108	37.670	1.00 32.44	AAAA
ATOM	731	CE	TY	2 Δ	91	23.250	38.203	38.475	1.00 31.88	AAAA
ATOM	732		YYT S			22.505	35.663	39.254	1.00 32.63	AAAA
ATOM	733	CE2	YYT S	٦Х	91	22.215	36.754	40.068	1.00 32.60	AAAA
ATOM	734	CZ	TYF	R A	91	22.589	38.016	39.668	1.00 31.59	AAAA
	735	OH	TYF			22.283				
ATOM							39.094	40.450	1.00 31.94	AAAA
MOTA	736	С	TYF			25.384	34.202	38.753	1.00 31.56	AAAA
ATOM	737	0	TYF	A S	91	26.075	35.105	39.233	1.00 31.21	AAAA
ATOM	738	N	GLU	TΑ	92	24.925	33.158	39.438	1.00 31.51	AAAA
	739	CA	GLU		92	25.143				
ATOM							32.941	40.865	1.00 32.70	AAAA
MOTA	740	CB	GLU	JA	92	24.463	31.626	41.268	1.00 33.55	AAAA
ATOM	741	CG	GLU	JA	92	24.174	31.495	42.747	1.00 34.16	AAAA
ATOM	742	CD	GLU	I A	92	23.311	30.278	43.087	1.00 35.31	AAAA
	743									
MOTA			GLU		92	23.857	29.148	43.152	1.00 34.30	AAAA
ATOM	744	OE2	GLU	A	92	22.076	30.466	43.275	1.00 35.36	AAAA
ATOM	745	С	GLU	I A	92	26.619	32.902	41.248	1.00 33.02	AAAA
ATOM	746	0	GLU	r A	92	27.073	33.623	42.140	1.00 32.91	AAAA
	747				93					
ATOM		N	ASN			27.358	32.049	40.550	1.00 32.84	AAAA
ATOM	748	CA	ASN	A	93	28.785	31.861	40.777	1.00 31.92	AAAA
MOTA	749	CB	ASN	I A	93	29.015	30.437	41.278	1.00 31.18	AAAA
ATOM	750	CG	ASN	Δ 1	93	27.948	29.994	42.259	1.00 30.34	AAAA
	751		ASN			27.723		43.271		
ATOM					93		30.642		1.00 31.20	AAAA
ATOM	752	ND2	ASN	A	93	27.284	28.892	41.955	1.00 29.02	AAAA
MOTA	753	С	ASN	A	93	29.442	32.052	39.411	1.00 30.84	AAAA
ATOM	754	0	ASN	ΙΑ	93	29.823	31.082	38.758	1.00 30.82	AAAA
	755	N	PRO		94	29.605	33.309	38.975	1.00 29.56	
ATOM										AAAA
MOTA	756	CD	PRO		94	29.312	34.590	39.626	1.00 29.03	AAAA
MOTA	757	CA	PRO	A	94	30.209	33.564	37.671	1.00 28.89	AAAA
ATOM	758	CB	PRO	Α	94	29.890	35.045	37.416	1.00 28.22	AAAA
ATOM	759	CG	PRO		94	28.839	35.377	38.435	1.00 29.50	AAAA
MOTA	760	C	PRO			. 31.698	33.351	37.664	1.00 28.25	AAAA
ATOM	761	Ø	PRO	A	94	32.308	32.996	38.671	1.00 28.21	AAAA
MOTA	762	N	VAL	A	95	32.257	33.593	36.488	1.00 27.36	AAAA
ATOM	763	CA	VAL		95	33.676	33.530	36.247	1.00 26.24	AAAA
ATOM	764	CB	VAL		95	33.945	33.289	34.741	1.00 26.10	AAAA
MOTA	765	CGI	VAL	Α	95	35.373	33.717	34.357	1.00 25.47	AAAA
ATOM	766	CG2	VAL	Α	95	33.736	31.826	34.434	1.00 25.59	AAAA
ATOM	767	С	VAL	A	95	34.178	34.919	36.647	1.00 26.56	AAAA
	768	ō	VAL		95	33.560	35.937	36.307	1.00 27.18	
ATOM										AAAA
MOTA	769	u	SER		96	35.280	34.966	37.382	1.00 25.23	AAAA
ATOM	770	CA	SER	Α	96	35.858	36.237	37.790	1.00 24.51	AAAA
ATOM	771	CB	SER	Α	96	34.935	36.961	38.774	1.00 23.22	AAAA
ATOM	772	OG	SER		96	34.941	36.297	40.014	1.00 19.76	AAAA
ATOM	773	С	SER		96	37.169	35.920	38.485	1.00 24.84	AAAA
MOTA	774	0	SER	A	96	37.590	34.764	38.530	1.00 25.97	AAAA
ATOM	775	N	TYR	Α	97	37.824	36.933	39.030	1.00 24.02	AAAA
ATOM	776	CA	TYR		97	39.047	36.664	39.744	1.00 24.55	AAAA
MOTA	777	CB	TYR		97	40.071	37.762		1.00 23.94	AAAA
MOTA	778	CG	TYR	A	97	40.682	37.636		1.00 23.72	AAAA
ATOM	779	CD1	TYR	A	97	40.177	38.341	37.039	1.00 23.11	AAAA
	780		TYR		97	40.700	38.136	35.758	1.00 23.50	AAAA
ATOM										
MOTA	781		TYR		97	41.717	36.735	37.903	1.00 22.25	AAAA
ATOM	782	CE2	TYR		97	42.236	36.526	36.640	1.00 22.86 ·	AAAA
ATOM	783	CZ	TYR	A	97	41.730	37.217	35.572	1.00 23.56	AAAA
	784	OH	TYR		97	42.232	36.941	34.318	1.00 24.06	AAAA
ATOM										
MOTA	785	<u>.</u> C	TYR		97	38.800	36.436	41.228	1.00 25.08	AAAA
ATOM	786	0	TYR	A	97	39.739	36.266	42.009	1.00 26.91	AAAA
ATOM	787	N	ALA		98	37.522	36.406	41.589	1.00 24.73	AAAA
		CA	ALA		98		36.159	42.951	1.00 24.50	AAAA
ATOM	788					37.083				
MOTA	789	CB	ALA		98	35.800	36.925	43.235	1.00 24.48	AAAA
MOTA	790	С	ALA	A	98	36.824	34.661	43.088	1.00 23.95	AAAA
ATOM	791	ο .	ALA	A	98	36.929	34.100	44.171	1.00 24.21	AAAA
•			MET		99			41.976	1.00 23.10	
ATCM	792	N	LIG I	u	2 <b>9</b>	36.502	34.011		1,00 43.10	AAAA
•										•

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ATOM	793 CA MET A 99	36.208	32.584	42.000	1.00 22.61	AAAA
MOTA	794 CB MET A 99	35.855				
ATOM	795 CG MET A 99	37.009				· AAAA
ATOM						AAAA
		36.360		•		AAAA
ATOM	797 CE MET A 99	35.328				AAAA
ATOM	798 C MET A 99	37.319	31.720	42.581	1.00 21.80	AAAA
MOTA	799 O MET A 99	37.052	30.695	43.199		AAAA
ATOM	800 N PHE A 100	38.567				
ATOM	801 CA PHE A 100					AAAA
		39.650				AAAA
ATOM	802 CB PHE A 100	40.388			1.00 20.25	AAAA
ATOM	803 CG PHE A 100 .	. 41.451			1.00 20.14	AAAA
ATOM	804 CD1 PHE A 100	41.114	28.462	43.010		AAAA
MOTA	805 CD2 PHE A 100	42.785			1.00 19.82	•
ATOM	806 CE1 PHE A 100	42.090				AAAA
	807 CE2 PHE A 100		27.695		1.00 19.54	AAAA
ATOM		43.755	29.300		1.00 19.22	AAAA
ATOM	808 CZ PHE A 100	43.410	28.122	43.641	1.00 19.47	AAAA
ATOM	809 C PHE A 100	40.649	32.161	43.743	1.00 21.37	AAAA
ATOM	810 O PHE A 100	40.959	31.822	44.887	1.00 21.26	AAAA
ATOM	811 N THR A 101	41.142	33.252		1.00 20.94	
ATOM	812 CA THR A 101					AAAA
		42.119	34.097		1.00 21.95	AAAA
ATOM	813 CB THR A 101	42.691	35.181		1.00 22.21	AAAA
ATOM	814 OG1 THR A 101	43.511	34.552	41.917	1.00 22.90	AAAA
ATOM	815 CG2 THR A 101	43.535	36.186		1.00 21.38	AAAA
ATOM	816 C THR A 101	41.584	34.755	45.117	1.00 22.60	AAAA
ATOM	817 O THR A 101	42.248	34.723		1.00 23.38	
ATOM	818 N GLY A 102					AAAA
		40.394	35.343		1.00 22.13	AAAA
ATOM	819 CA GLY A 102	39.826	35.972	46.227	1.00 22.03	AAAA
MOTA	820 C GLY A 102	39.340	34.928	47.221	1.00 21.36	AAAA
MOTA	821 O GLY A 102	39.433	35.104	48.439	1.00 20.02	AAAA
MOTA	822 N SER A 103	38.816	33.833	46.677	1.00 21.86	AAAA
ATOM	823 CA SER A 103	38.311	32.719	47.466	1.00 21.68	
ATOM	824 CB SER A 103	37.699	31.668	46.557		AAAA
ATOM					1.00 21.56	AAAA
		36.604	32.216	45.857	1.00 23.67	AAAA
ATOM	826 C SER A 103	39.450	32.098	48.229	1.00 22.67	AAAA
MOTA	827 O SER A 103	39.314	31.806	49.412	1.00 22.44	AAAA
ATOM	828 N SER A 104	40.578	31.898	47.545	1.00 23.37	AAAA
ATOM	829 CA SER A 104	41.746	31.305	48.183	1.00 23.50	AAAA
ATOM	830 CB SER A 104	42.862	31.070	47.172	1.00 24.80	AAAA
ATOM	831 OG SER A 104	42.441	30.169	46.175		
ATOM					1.00 28.38	AAAA
		42.254	32.230	49.256	1.00 22.79	AAAA
ATOM	833 O SER A 104	42.707	31.794	50.307	1.00 22.66	AAAA
ATOM	834 N LEU A 105	42.160	33.518	48.970	1.00 22.08	AAAA
MOTA	835 CA LEU A 105	42.626	34.541	49.870	1.00 21.70	AAAA
ATOM	836 CB LEU A 105	42.524	35.882	49.159	1.00 21.89	AAAA
MOTA	837 CG LEU A 105 ·	43.332	37.038	49.718	1.00 23.64	
ATOM	838 CD1 LEU A 105	44.830				AAAA
			36.692	49.639	1.00 22.01	AAAA
MOTA	839 CD2 LEU A 105	43.004		48.919	1.00 23.60	AAAA
ATOM	840 C LEU A 105	41.767	34.525	51.131	1.00 22.29	AAAA
MOTA	841 O LEU A 105	42.277	34.595	52.249	1.00 21.95	AAAA
ATOM	842 N ALA A 106	40.458	34.429	50.934	1.00 22.23	AAAA
MOTA	843 CA ALA A 106	39.515		52.042	1.00 22.32	AAAA
ATOM	844 CB ALA A 106	38.068	34.472		1.00 22.05	
				51.526		AAAA
ATOM	845 C ALA A 106	39.704	33.126	52.840	1.00 21.99	AAAA
ATOM	846 O ALA A 106	39.5 <b>7</b> 8	33.145	54.061	1.00 23.18	AAAA
ATOM	847 N THR A 107	40.011	32.032	52.144	1.00 21.24	AAAA
MOTA	848 CA THR A 107	40.209	30.732	52.779	1.00 20.60	AAAA
ATOM	849 CB THR A 107	40.170	29.571	51.749	1.00 19.82	
MOTA	850 CG1 THR A 107					AAAA
		38.903	29.553	51.083	1.00 18.56	AAAA
ATOM	851 CG2 THR A 107	40.360	28.242	52.455	1.00 18.58	Aaaa
MOTA	852 C THR A 107.	41.516	30.630	53.561	1.00 21.41	AAAA
ATOM	853 O THR A 107	41.537	30.040	54.646	1.00 23.16	AAAA
ATOM	854 N GLY A 108	42.601	31.176	53.003	1.00 20.14	AAAA
ATOM	855 CA GLY A 108	43.878	31.145	53.684	1.00 18.20	
						AAAA
ATOM .	856 C GLY A 108	43.739	31.933	54.972	1.00 18.43	AAAA
ATOM	857 O GLY A 108	44.335	31.600	55.998	1.00 17.52	AAAA
ATOM	858 N SER A 109	42.909	32.969	54.929	1.00 18.56	. AAAA

ATOM	859	CA	SER	A 109	42.683	33.805	56.098	1.00 19.67	AAAA
	860	CB		A 109	41.899	35.058	55.707	1.00 20.27	AAAA
ATOM .									
ATOM	861	OG	SER	A 109	42.618	35.803	54.746	1.00 21.80	AAAA
MOTA	862	С	SER	A 109	41.955	33.066	57.219	1.00 19.61	AAAA
•	863	ō		A 109	42.078	33.426	58.388	1.00 18.40	AAAA
ATOM		-							
ATOM -	864	N	THR	A 110	41.186	32.042	56.866	1.00 19.88	AAAA
MOTA	865	CA	THR	A 110	40.493	31.288	57.891	1.00 20.51	AAAA
						30.438	57.304	1.00 20.62	AAAA
MOTA	866	CB		A 110	39.365				
· ATOM ·	867	OG1	THR	A 110	38.236	31.284	57.050	1.00 20.80	AAAA
	868	CG2	THR	A 110	38.974	29.313	58.262	1.00 20.53	AAAA
ATOM							58.601	1.00 20.36	AAAA
MOTA	869	С		A 110	41.504	30.420			
MOTA	870	0	THR	A -110	41.455	30.268	59.822	1.00 20.78	AAAA
	871	N	VAT.	A 111	42.431	29.855	57.832	1.00 20.85	AAAA
ATOM							58.423	1.00 21.03	AAAA
ATOM	872	CA		A 111	43.480	29.053		· ·	
MOTA	873	CB	VAL	A 111	44.318	28.323	57.345	1.00 21.05	AAAA
	874	CGI	VAT.	A 111	45.537	27.644	57.983	1.00 19.91	-AAAA
MOTA						27.281	56.648	1.00 18.39	AAAA
MOTA	875			A 111	43.460				
ATOM	876	С	VAL	A 111	44.374	30.005	59.232	1.00 21.84	AAAA
ATOM	877	0	WAT.	A 111	44.825	29.671	60.331	1.00 22.73	· AAAA
						31.204	58.712		AAAA
ATOM	878	N		A 112	44.612				
ATOM	879	CA	GLN .	A 112	45.449	32.133	59.452	1.00 21.89	AAAA
ATOM	880	CB	GLN	A 112	45.630	33.450	58.6 <del>9</del> 0	1.00 22.50	AAAA
	881			A 112	46,288	33.283	57.335	1.00 23.68	AAAA
ATOM		CG							
ATOM	882	CD	GLN	A 112	46.414	34.578	56.569	1.00 23.18	AAAA
MOTA	883	OE1	GLN	A 112	47.389	35.310	56.722	1.00 23.86	AAAA
	884			A 112	45.413	34.879	55.752	1.00 21.90	AAAA
MOTA									
ATCM	885	С	GLN	A 112	44.766	32.383	60.774	1.00 21.84	AAAA
ATOM	886	0	GLN .	A 112	45.389	32.316	61.835	1.00 22.47	AAAA
	887	N		A 113	43.468	32.651	60.700	1.00 21.34	AAAA
MOTA					42.500		61.884	1.00 20.84	AAAA
MOTA	888	CA		A 113	42.682	32.934			
ATOM	889	CB	ALA	A 113	41.244	33.172	61.504	1.00 18.52	AAAA
ATOM	890	С	A.T.A	A 113	42.795	31.782	62.865	1.00 21.75	AAAA
					42.880	31.985	64.084	1.00 22.24	· AAAA
MOTA	891	0		A 113					
ATOM	892	N	ILE .	A 114	42.797	30.569	62.329	1.00 22.54	AAAA
MOTA	893	CA.	ILE	A 114	42.891	29.393	63.160	1.00 23.16	AAAA
	894	CB		A 114	42.557	28.146	62.352	1.00 23.33	AAAA
ATOM							63.106	1.00 23.80	AAAA
ATOM	895			A 114	42.939	26.912			
MOTA	896	CG1	ILE	A 114	41.058	28.130	62.047	1.00 23.48	AAAA
	897	CD1	TLE	A 114	40.610	26.951	61.204	1.00 22.08	AAAA
MOTA						29.270	63.792	1.00 24.33	AAAA
ATOM	898	С		A 114	44.268			•	
ATOM	899	0	ILE .	A 114	44.373	29.013	64.990	1.00 25.30	AAAA
ATOM	900	N	GLU	A 115	45.319	29.490	63.002	1.00 24.96	AAAA
	_	CA		A 115	46.699	29.395	63.503	1.00 26.61	AAAA
ATOM	901							1.00 24.75	AAAA
ATOM	902	CB	GLU .	A 115	47.708	29.753	62.406		
ATOM	903	CG	GLU .	A 115	47.444	29.033	61.103	1.00 25.80	AAAA
	904	CD		A 115	48.471	29.323	60.030	1.00 26.07	AAAA
MOTA							59.940	1.00 27.15	AAAA
ATOM	905			A 115	48.911	30.484			
ATOM	906	OE2	GLU .	A 115	48.819	28.402	59.260	1.00 25.45	AAAA
ATOM	907	С	GLU	A 115	46.877	30.340	64.680	1.00 27.89	AAAA
		ŏ		A 115	47.480	29.975	65.695	1.00 28.04	AAAA
ATOM	908					31.552	64.531	1.00 29.15	AAAA
ATOM	909	N	GLU .	A 116	46.337	31.352			
MOTA	910	CA	GLU	A 116	46.408	32.579	65.563	1.00 29.42	AAAA
					45.751	33.871	65.082	1.00 28.26	AAAA
ATOM	911	CB		A 116				1.00 28.93	AAAA
ATOM	912	CG	GLU .	A 116		34.529	63.945		
ATOM	913	CD	GLU .	A 116	47.902	34.937	64.318	1.00 28.32	AAAA
	914			A 116	48.081	35.878	65.123	1.00 27.68	AAAA
ATOM							63.810	1.00 27.38	AAAA
ATOM	91,5	OE2	GLU .	A 116	48.838	34.297			
ATOM	916	С	GLU .	A 116	45.737	32.126	66.845	1.00 29.77	AAAA
		ō		A 116	46.338	32.196	67.920	1.00 30.29	AAAA
ATOM	917						66.727	1.00 29.64	
ATOM	918	N		A 117	44.492	31.665			
ATOM	919	CA	PHE	A 117	43.741	31.204	67.88 <b>7</b>	1.00 29.33	AAAA
	920	CB		A 117	42.425	30.552	67.480	1.00 28.89	AAAA
ATOM						30.087	68.651	1.00 28.93	AAAA
ATCM	921	CG		A 117	41.604				
ATOM	922	CD1	PHE .	A 117	41.010	31.010	69.510	1.00 28.42	AAAA
	923			A 117	41.441	28.723	68.910	1.00 29.06	AAAA
ATCM				A 117	40.261	30.588	70.610	1.00 28.68	AAAA
	924	للظا	PME .	W TT/	40.201	30.300			*******
ATOM	204								

ATOM	925	CE2	. PHE A	A 117	40.695	28.284	70.009	1.00 29.1	т аааа
	926	cz			40.103	29.227	70.862	1.00 29.0	
MOTA				A 117					
ATOM	927	С	PHE A	A 117	44.545	30.195	68.671	1.00 29.2	22 AAAA
ATOM	928	0	PHE A	A 117	44.677	30.315	69.884	1.00 30.2	29 AAAA
	929	N		A 118	45.066	29.195	67.965	1.00 29.2	
ATOM									
ATOM	930	CA	LEU A	A 118	45.864	28.145	68.576	1.00 29.5	AAAA 0
ATOM	931	CB	LEUR	4 118	46.182	27.047	67.550	1.00 28.5	57 AAAA
				118		26.296	66.989	1.00 28.1	
ATOM	932	CG			44.962				
ATOM	933	CD1	LEU A	A 118	45.421	25.090	66.191	1.00 25.5	58 AAAA
ATOM	934	CD2	LEU 2	118	44.053	25.846	68.128	1.00 27.6	54 AAAA
						28.649	69.227	1.00 30.1	
ATOM	935	C	LEU A		47.150				
ATOM	936	0	LEU A	1118	47.727	27.954	70.056	1.00 29.9	94 AAAA
ATOM	937	N	LYS A	119	47.602	29.845	68.847	1.00 31.3	BG AAAA `
	938	CA	LYS A		48.798	30.451	69.448	1.00 32.5	
MOTA									
MOTA	939	CB	LYS A	119	49.396	31.539	68.559	1.00 32.3	
ATOM	940	CG	LYS A	119	49.882	31.108	67.199	1.00 33.0	)3 AAAA
	941	CD	LYS A		50.371	32.321	66.411	1.00 32.7	
ATOM								•	
ATOM	942	CE	LYS A	1119	50.681	31.939	64.972	1.00 33.9	
MOTA	943	NZ	LYS A	119	51.125	33:099	64.152	1.00 34.9	эз аааа
	944	С	LYS A		48.385	31.143	70.744	1.00 33.7	4 AAAA
ATOM									
ATOM	945	0	LYS A	1119	49.218	31.748	71.413	1.00 34.8	
MOTA	946	N	GLY A	120	47.096	31.079	71.073	1.00 33.6	AAAA 88
	947	CA	GLY A		46.600	31.736	72.263	1.00 33.6	59 AAAA
MOTA								1.00 34.1	
ATOM	948	С	GLY A		45.987	33.110	71.988		
ATOM	949	0	GLY A	120	45.588	33.802	72.932	1.00 33.6	55 AAAA
MOTA	950	N	ASN A	121	45.904	33.513	70.717	1.00 33.5	AAAA 8
	951	CA	ASN A		45.326	34.820	70.368	1.00 33.3	
MOTA									
ATOM	952	CB	ASN A	121 .	46.194	35.537	69.341	1.00 33.1	
ATOM	953	CG	ASN A	121	47.570	35.828	69.859	1.00 34.3	31 AAAA
ATOM	954	വാ	ASN A	121	48.333	34.921	70.154	1.00 35.6	57 AAAA
					47.897	37.096	69.975	1.00 34.1	
MOTA	955		ASN A						
MOTA	956	С	ASN A	. 121	43.888	34.805	69.839	1.00 32.8	
MOTA	957	0	ASN A	121	43.304	33.751	69.599	1.00 32.7	78 AAAA
	958	N	VAL A		43.338	36.003	69.655	1.00 32.4	7 AAAA
MOTA									
ATOM	959	CA	VAL A		41.980	36.200	69.148	1.00 30.8	
ATOM	960	CB	VAL A	122	41.182	37.145	70.070	1.00 31.0	)5 AAAA
ATOM	961	CGI	VAL A	122	39.831	37.423	69.489	1.00 30.9	AAAA C
					41.038	36.516	71.440	1.00 31.1	
MOTA	962		VAL A						
MOTA	963	С	VAL A	122	42.056	36.805	67.750	1.00 30.1	
ATOM .	964	0	VAL A	. 122	42.694	37.840	67.535	1.00 31.2	AAAA 8
	965	N	ALA A		41.405	36.147	66.800	1.00 28.6	52 AAAA
ATOM							65.421	1.00 26.4	
MOTA	966	CA	ALA A		41.415	36.589			
ATOM	967	CB	ALA A	. 123	42.323	35.708	64.599	1.00 26.5	
ATOM	968	С	ALA A	123	40.038	36.570	64.836	1.00 25.5	59 AAAA
	969	ō	ALA A		39.173	35.814	65.252	1.00 26.2	27 AAAA
ATOM									
ATOM	970	N	PHE A		39.848	37.421	62.847	1.00 25.4	
ATOM	971	CA	PHE A	. 124	38.590	37.534	61.156	1.00 23.8	37 AAAA
ATOM	972	CB	PHE A	124	37.832	38.779	62.646	1.00 23.5	SS AAAA
					36.591	39.119	62.841	1.00 23.7	1 AAAA
MOTA	973	CG	PHE A					•	
ATOM.	974	CD1	PHE A	124	35.668	38.140	62.495	1.00 23.4	
MOTA	975	CD2	PHE A	124	36.311	40.449	62.498	1.00 23.7	
	976		PHE A		34.479	38.483	61.823	1.00 23.3	AAAA
ATOM							61.833	1.00 21.7	
MOTA	977	CEZ	PHE A		35.131	40.796			
ATOM .	978	CZ	PHE A	124	34.217	39.815	61.497	1.00 22.3	35 AAAA
ATOM	979	С	PHE A	124	38.951	37.673	61.700	1.00 23.2	26 AAAA
			PHE A		39.720	38.555	61.323	1.00 22.2	
ATOM	980	0						1.00 23.2	
ATOM	981	N	ASN A		38.427	36.759	60.897		
ATOM	982	CA	ASN A	125	38.622	36.785	59.457	1.00 21.0	
		СВ	ASN A		39.181	35.470	58.951	-1.00 19.9	
ATOM	983						57.454	1.00 20.6	
ATOM	984	CG	ASN A		39.098	35.360			
MOTA	285	OD1	ASN A	125	39.389	36.317	56.748	1.00 21.6	
	986		ASN A		38.721	34.190	56.956	1.00 19.9	AAAA E
ATOM			3 (2)7 3	125	37.269	37.059	58.813	1.00 20.1	
MOTA	987	C	ASN A	105					
ATOM	988		ASN A		36.469	36.148	58.579	1.00 19.2	
ATOM	989	N	PRO A	126	36.991	38.340	58.543	1.00 19.1	
•		CD	PRO A	126	37.893	39.460	58.858	1.00 19.2	2 AAAA
ATOM	990	CD,	T.UO M		3	22.200			

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ATOM	991	L CA	PRO A 12	5 35.766	38.849	57.932	1.00 19.52	AAAA
ATOM	992	2 CB	PRO A 120	36.005				
ATOM	993	CG						AAAA
ATOM	994		PRO A 126					AAAA
ATOM	. 995						1.00 19.41	AAAA
	-		PRO A 126				1.00 19.68	AAAA
MOTA	996		ALA A 127			55.835	1.00 18.17	AAAA
ATOM	997	' CA	ALA A 127	36.283	37.314	54.481	1.00 17.66	AAAA
ATOM	998	CB	ALA A 127	37.547	37.520		1.00 17.08	-
ATOM	999	C	ALA A. 127				1.00 17.46	AAAA
ATOM	1000		ALA A 127				1.00 17.46	AAAA
ATOM	1001		GLY A 128				1.00 18.92	AAAA
							1.00 15.94	AAAA
MOTA	1002						1.00 15.45	AAAA
MOTA	1003		GLY A 128	34.226	33.593	55.955	1.00 16.08	AAAA
ATOM	1004	0	GLY A 128	33.485	34.557	55.997	1.00 15.43	AAAA
ATOM	1005	N	GLY A 129	33.821	32.353		1.00 16.77	AAAA
ATOM	1006	CA	GLY A 129		32.082		1.00 17.82	
ATOM	. 1007		GLY A 129		31.822			AAAA
ATOM	1008		GLY A 129	_			1.00 18.64	AAAA
					32.051		1.00 18.48	AAAA
MOTA	1009		MET A 130		31.368		1.00 20.45	AAAA
ATOM	1010		MET A 130		31.029	52.826	1.00 21.60	AAAA
MOTA	1011	CB	MET A 130		31.117	51.744	1.00 22.02	AAAA
ATOM	1012	CG	MET A 130	33.551	32.472		1.00 21.75	AAAA
ATOM	1013	SD	MET A 130	34.971	32.567	50.599	1.00 24.75	
ATOM	1014	CE	MET A 130		32.137	49.048		AAAA
ATOM	1015	c	MET A 130	31.328			1.00 24.40	AAAA
ATOM	1016	õ			29.587	53.002	1.00 22.08	AAAA
			MET A 130	31.970	28.641	52.546	1.00 22.98	AAAA
ATOM	1017	N	HIS A 131	30.184	29.452	53.659	1.00 22.25	AAAA
ATOM	1018	CA	HIS A 131	29.618	28.171	54.062	1.00 20.49	AAAA
ATOM	1019	CB	HIS A 131	28.832	28.421	55.342	1.00 20.00	AAAA
ATOM	1020	CG	HIS A 131	27.679	29.360	55.161	1.00 17.93	AAAA
ATOM	1021	CD2	HIS A 131	27.091	29.846	54.043	1.00 17.88	AAAA
ATOM	1022		HIS A 131	26.952	29.854	56.219	1.00 19.33	•
MOTA	1023		HIS A 131	25.968	30.607			AAAA
ATOM	1024		HIS A 131			55.758	1.00 16.99	AAAA
				26.031	30.617	54.441	1.00 17.43	AAAA
ATOM	1025	C	HIS A 131	28.763	27.332	53.141	1.00 19.97	AAAA
ATOM	1026	0	HIS A 131	28.330	26.262	53.541	1.00 19.61	· AAAA
ATOM	1027	N	HIS A 132	28.518	27.796	51.923	1.00 20.11	AAAA
ATOM	1028	CA	HIS A 132	27.673	27.058	50.994	1.00 17.76	AAAA
ATOM	1029	CB	HIS A 132	26.879	28.044	50.127	1.00 16.76	AAAA
ATOM	1030	CG	HIS A 132	25.824	28.815	50.862	1.00 15.35	AAAA
ATOM	1031	CD2	HIS A 132	25.567	30.146	50.920	1.00 14.15	
ATOM	1032		HIS A 132	24.804	28.200			AAAA
ATOM	1033		HIS A 132			51.557	1.00 16.15	AAAA
	1034			23.966	29.119	52.005	1.00 14.13	AAAA
ATOM			HIS A 132	24.405	30.307	51.632	1.00 14.65	AAAA
ATOM	1035	C	HIS A 132	28.355	26.051	50.065	1.00 17.99	AAAA
MOTA	1.36	0	HIS A 132	27.742	25.053	49.684	1.00 18.54	- AAAA
ATOM	1:J37	N	ALA A 133	29.604	26.305	49.690	1.00 17.82	AAAA
ATOM	1638	CA	ALA A 133	30.300	25.441	48.742	1.00 18.38	AAAA
ATOM	1039	CB	ALA A 133	31.684	25.961	48.507	1.00 17.53	AAAA
ATOM	1040	С	ALA A 133	30.366	23.970	49.130	1.00 20.92	
ATOM	1041	ō	ALA A 133	30.578				AAAA
				_	23.633	50.298	1.00 21.79	AAAA
ATOM	1042	N	PHE A 134	30.184	23.086	48.152	1.00 20.58	AAAA
ATOM	1043		PHE A 134	30.258	21.663	48.455	1.00 21.38	AAAA
ATOM	1044		PHE A 134	29.168	20.860	47.731	1.00 19.41	AAAA .
MOTA	1045	CG	PHE A 134	27.772	21.229	48.126	1.00 18.32	AAAA
ATOM	1046	CD1	PHE A 134	27.027	22.099	47.357	1.00 19.22	AAAA
ATOM	1047		PHE A 134	27.193	20.701	49.271	1.00 19.14	
ATOM	1048		PHE A 134	25.714		47.726		AAAA
ATOM					22.438		1.00 18.56	AAAA
	1049		PHE A 134	25.889	21.036	49.644	1.00 17.72	AAAA
ATOM	1050		PHE A 134	25.158	21.903	48.866	1.00 18.01	AAAA
MOTA	1051		PHE A 134	31.625	21.124	48.081	1.00 22.90	AAAA
MOTA	1052	0	PHE A 134	32.459	21.833	47.544	1.00 23.37	AAAA
MOTA	1053		LYS A 135	31.842	19.861	48.390	1.00 24.63	AAAA
ATOM	1054		LYS A .135	33.095	19.195	48.122	1.00 27.16	
ATOM	1055		LYS A 135	32.926			1.00 28.53	AAAA
					17.714	48.480		AAAA
ATOM	1056	CG	LYS A 135	34.133	16.843	48.292	1.00 31.01	AAAA

ATOM	1057	CD	LYS A 13	33.879	15.472	48.910	1.00 32.75	AAAA
MOTA	1058	CE	LYS A 13	33.961	15.495	50.457	1.00 33.96	AAAA
ATOM	1059	NZ	LYS A 13	35.371	15.664	50.976	1.00 33.04	AAAA
MOTA	1060	С	LYS A 13	33.577	19.390	46.673	1.00 27.37	AAAA
ATOM	1061	0	LYS A 13.	34.769	19.596	46.437	1.00 27.35	AAAA
ATOM	1062	N	SER A 13	32.658	19.354	45.714	1.00 27.32	AAAA
MOTA	1063	CA	SER A 13	33.028	19.527	44.313	1.00 28.31	AAAA
ATOM	1064	CB	SER A 13	33.093	18.162	43.626	1.00 28.56	AAAA
ATOM	1065	OG	SER A 13	33.822	17.242	44.417	1.00 29.28	AAAA
MOTA	1066	С	SER A 13	31.993	20.395	43.599	1.00 28.91	AAAA
ATOM	1067	0	SER A 13		20.080	42.486	1.00 28.78	AAAA
ATOM	1068	N	ARG A 13'		21.502	44.212	1.00 29.08	AAAA
ATOM	1069	CA	ARG A 13'		22.311	43.576	1.00 29.66	AAAA
ATOM	1070	CB	ARG A 13'	29.259	21.528	43.657	1.00 31.65	AAAA
ATOM	1071	CG	ARG A 13	27.989	22.273	43.355	1.00 33.89	AAAA
ATOM	1072	CD	ARG A 13	26.862	21.267	43.373	1.00 35.93	AAAA
ATOM	1073	NE	ARG A 13	26.961	20.366	42.228	1.00 36.31	AAAA
ATOM	1074	CZ	ARG A 137	26.505	20.660	41.015	1.00 35.99	AAAA
MOTA	1075	NH1	ARG A 137		21.834	40.798	1.00 34.63	AAAA
ATOM	1076	NH2	ARG A 137	26.650	19.786	40.025	1.00 35.35	AAAA
ATOM	1077	·C	ARG A 137	30.402	23.723	44.116	1.00 28.53	AAAA
MOTA	1078	0	ARG A 137	30.418	23.946	45.324	1.00 28.51	AAAA
ATOM	1079	N	ALA A 138	30.247	24.673	43.202	1.00 27.53	AAAA
ATOM:	1080	CA	ALA A 138	30.039	26.063	43.581	1.00 27.64	AAAA
MOTA	1081	CB	ALA A 138	30.236	26.984	42.381	1.00 27.87	AAAA
ATOM	1082	C	ALA A 138	28.601	26.130	44.079	1.00 27.27	AAAA
ATOM	1083	0	ALA A 138	27.769	25.321	43.671	1.00 28.30	AAAA
ATOM	1084	И	ASN A 139	28.292	27.080	44.951	1.00 26.16	AAAA
ATOM	1085	CA	ASN A 139	26.945	27.134	45.480	1.00 25.39	AAAA
ATOM	1086	CB	ASN A 139	26.673	25.847	46.282	1.00 24.58	AAAA
ATOM	1087	CG	ASN A 139	25.343	25.872	47.017	1.00 25.37	AAAA
ATOM	1088		ASN A 139	24.272	26.017	46.413	1.00 24.20	AAAA
ATOM	1089		ASN A 139	25.408	.25.720	48.338	1.00 24.91	AAAA
ATOM	1090	Ċ	ASN A 139		28.358	46.341	1.00 24.90	AAAA
ATOM	1091	0	ASN A 139	27.346	28.570	47.348	1.00 24.98	AAAA
MOTA	1092	N	GLY A 140	25.702	29.145	45.916	1.00 24.46	AAAA
MOTA	1093	CA	GLY A 140	25.294	30.336	46.625	1.00 22.96	AAAA
MOTA	1094	C	GLY A 140	26.383	31.358	46.755	1.00 22.24	AAAA
MOTA	1095	0	GLY A 140	26.663	31.817	47.867	1.00 23.09	AAAA
ATOM	1096	N	PHE A 141	26.992	31.711	45.625	1.00 20.60	AAAA
MOTA	1097	CA	PHE A 141	28.075	32.700	45.572	1.00 19.43	AAAA
MOTA	1098	CB CG	PHE A 141	27.758 26.453	33.920 34.577	46.430	1.00 19.86	AAAA
MOTA	1099 1100		PHE A 141 PHE A 141	25.974	35.592	46.114 46.934	1.00 21.18 1.00 20.49	AAAA
ATOM			PHE A 141	25.723	34.218	44.985	1.00 20.43	AAAA AAAA
ATOM "	1101		PHE A 141	24.800	36.242	46.638	1.00 21.42	AAAA
ATOM ATOM	1103		PHE A 141	24.540	34.859	44.672	1.00 :1.76	AAAA
MOTA	1104	CZ	PHE A 141	24.072	35.881	45.499	1.00 3.05	AAAA
ATOM	1105	c	PHE A 141	29.396	32.132	46.069	1.00 18.68	AAAA
ATOM	1106	ō	PHE A 141	30.438		45.944	1.00 19.19	AAAA
ATOM	1107	N	CYS A 142	29.367	30.930	46.635	1.00 16.93	AAAA
ATOM	1108	CA	CYS A 142	30.594	30.332	47.150	1.00 16.80	AAAA ·
ATOM	1109	CB	CYS A 142	30.323	29.689	48.509	1.00 16.51	AAAA
ATOM	1110	SG	CYS A 142	29.524	30.826	49.617	1.00 15.01	AAAA
ATOM	1111	C	CYS A 142	31.227	29.315	46.221	1.00 16.45	AAAA
ATOM	1112	õ.	CYS A 142	30.533	28.565	45.556	1.00 15.32	AAAA
ATOM	1113	N	TYR A 143	32.558	29.311	46.190	1.00 18.39	AAAA
ATOM	1114	CA	TYR A 143	33.340	28.394	45.362	1.00 18.63	AAAA
ATOM	1115		TYR A 143	34.298	29.154	44.438	1.00 19.48	AAAA
ATOM	1116		TYR A 143	33.664	30.214	43.571	1.00 19.55	AAAA
				33.480	31.510	44.043	1.00 20.27	AAAA
	1117	CDT	TYR A 143					
ATÓM	1117 1118		TYR A 143	32.856	32.473	43.261		AAAA
atom atom	1118	CEI	TYR A 143	32.856	32.473	43.261	1.00 21.63	AAAA
atom atom atom	1118 1119	CE1 CD2	TYR A 143 TYR A 143				1.00 21.63 1.00 20.14	AAAA AAAA
atom atom	1118	CE1 CD2 CE2	TYR A 143	32.856 33.212	32.473 29.910	43.261 42.292	1.00 21.63	AAAA

ATOM 1121 CZ TIR A 143 31.787 33.071 4

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ATOM	1123	С	ጥሃፑ	A	143		34,162	27.490	46.283	1.00	19.06	AAAA
					143		34.319	26.289	46.032		18.40	AAAA
MOTA	1124	0										
ATOM	1125	N	ILE	A	144		34.695	28.087	47.344		19.15	AAAA
ATOM	1126	CA	ILE	Α	144		35.490	27.350	48.315	1.00	19.97	AAAA
	1127	CB			144		36.952	27.861	48.355		19.74	AAAA
ATOM												
ATOM	1128		ILE				37.757	27.088	49.410		18.03	AAAA
MOTA	1129	CG1	ILE	A	144		37.584	27.671	46.965	1.00	20.12	AAAA
ATOM	1130	CD1	ILE	Α	144		39.053	28.072	46.846	1.00	21.05	`AAAA
							34.833	27.532	49.665	-	20.22	AAAA
MOTĄ	1131	C			144							
ATOM	1132	0	LILE	A	144		34.357	28.626	49.981	1.00	19.94	AAAA
MOTA	1133	N	ASN	Α	145		34.787	26.451	50.440	1.00	20.57	AAAA
	1134	CA	ASN				34.165	26.448	51.770	1 00	20.39	AAAA
ATOM									51.990			
MOTA	1135	CB			145		33.450	25.114			19.39	AAAA
MOTA	1136	CG	ASN	Α	145		32.505	25.143	53.171	1.00	19.31	AAAA
ATOM	1137	OD1	ASN	Α	145		32.862	25.583	54.263	1.00	21.26	AAAA
	1138		ASN				31.290	24.667	52.960		17.08	- AAAA
ATOM												
ATOM	1139	С			145		35.236	26.621	52.856		20.17	AAAA
ATOM	1140	0	ASN	Α	145		35.690	25.622	53.421	1.00	19.75	AAAA
ATOM	1141	N	ASN	Α	146	•	35.644	27.862	53.148	1.00	20.06	AAAA
		CA	ASN				36.671	28.075	54.166		20.98	AAAA
ATOM	1142											
ATOM	1143	CB	ASN				37.019	29.573	54.333		21.78	AAAA
ATOM	1144	CG	ASN	Α	146		35.876	30.411	54.882	1.00	22.78	AAAA
ATOM	1145	OD3	ASN	А	146		35.651	30.465	56.091	1.00	22.83	AAAA
			ASN				35.144	31.078	53.983		23.70	AAAA
ATOM	1146											
ATOM ·	1147	C	asn	Ą	146		36.307	27.413	55.496		21.18	AAAA
ATOM	1148	0	ASN	Α	146		37.169	26.823	56.139	1.00	21.48	AAAA
ATOM	1149	N	PRO	À	147		35.031	27.476	55.922	1.00	20.88	AAAA
	1150	CD	PRO				33.835	28.120	55.358		21.85	AAAA
ATOM												
MOTA	1151	CA	PRO				34.674	26.831	57.183		21.42	AAAA
ATOM	1152	CB	PRO	А	147		33.176	27.073	57.261	1.00	21.00	AAAA
ATOM	1153	CG	PRO	А	147	•	33.052	28.408	56.605	1.00	20.47	AAAA
	1154	C	PRO.		_		35.015	25.334	57.174		22.79	AAAA
ATOM												
ATOM	1155	0	PRO				35.650	24.833	58.099		25.69	AAAA
ATOM	1156	N	ALA	А	148		34.603	24:616	56.136	1.00	22.34	AAAA
ATOM	1157	CA	ALA	Α	148		34.889	23.193	56.070	1.00	22.23	AAAA
	1158	CB	ALA		_		34.260	22.561	54.825	1.00	22.87	AAAA
MOTA									56.054		22.33	AAAA
ATOM	1159	C.	ALA				36.378	22.998				
ATOM	1160	0	ALA	Α	148		36.912	22.249	56.861		23.42	AAAA
MOTA	1161	N	VAL	Α	149		37.050	23.661	55.122	1.00	22.50	AAAA
	1162	CA	VAL				38.505	23.569	55.018	1.00	21.29	AAAA
MOTA							39.066	24.581	54.002		20.46	AAAA
MOTA	1163	CB	VAL									
ATOM	1164	CG1	VAL	A	149		40.578	24.607	54.085		19.36	AAAA
ATOM	1165	CG2	VAL	Α	149		38.608	24.229	52.593	1.00	20.03	AAAA
MOTA	1166	С	VAL	Α	149		39.164	23.848	56.367	1.00	21.48	AAAA
		ŏ	VAL				40.147	23.197	56.735		22.11	AAAA
ATOM	1167										21.19	
MOTA	1168	N	GL.				38.628	24.826	57.088			AAAA
MOTA	1169	CA	GL.	А	150		39.171	25.176	58.386		21.70	. AAAA
MOTA	1170	С.	GL'_	А	150		38.973	24.043	59.368	1.00	22.31	AAAA
	1171	ō	GLY				39.913	23.597	60.026	1.00	22.51	AAAA
ATOM									59.453		22.86	AAAA
ATOM	1172	N	ILE				37.736	23.566				
ATOM	1173	CA	ILE	A	151		37.388	22.474	60.346		22.26	AAAA
ATOM	1174	CB	ILE	Α	151		35.894	22.124	60.191	1.00	21.51	AAAA
			ILE				35.542	20.899	61.019		21.36	AAAA
ATOM	1175								60.627		20.39	AAAA
ATOM	1176		ILE				35.051	23.329				
ATOM	1177	CD1	ILE	A	151		33.576	23.199	60.361		16.88	AAAA
ATOM	1178	С	ILE	Α	151		38.265	21.243	60.096	1.00	23.29	AAAA
			ILE				38.786	20.660	61.038		2388	AAAA
ATOM	1179	0										
ATOM	1180		GLU				38.435	20.853	58.836		24.13	AAAA
ATOM	1181	CA	GLU	Α	152		39.267	19.697	58.517	1.00	25.01	AAAA
	1182		GLU				39.242	19.404	57.010	1.00	25.07	AAAA
ATOM			GLU				37.910	_	56.526		24.56	AAAA
atom	1183							18.886			25.00	
ATOM	1184	CD	GLU	A	125		37.500	17.570	57.198			AAAA
ATOM	1185	OE1	GLU	Α	152		36.345	17.158	57.011		26.40	AAAA
	1186	OF2	GLU	А	152		38.315	16.935	57.897	1.00	25.00	AAAA
ATOM		~	GLU	Δ	152		40.694	19.957	58.965		26.06	AAAA
ATOM	1187	c	OTIO .	•	150						26.40	AAAA
MOTA	1188	0	GLU	A	122		41.425	19.035	59.331	1.00	40.40	, AAAA

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### Figure 19-19

ATOM	1189	N	TYR I	A 153	41.085	21.225	58.925	1.00 27.30	AAAA
MOTA	1190	CA	TYR	A 153	42.422	21.632	59.334	1.00 27.63	AAAA
	1191	CB		A 153	42.532	23.153	59.268	1.00 26.99	
ATOM					_				AAAA
MOTA	1192	CG		A 153	43.856	23.719	59.710	1.00 27.03	· AAAA
MOTA	1193	CD	l TYR I	A 153	44.942	23.790	58.837	1.00 27.78	AAAA
ATOM	1194	CE1	L TYR I	4 153	46.165	24.356	59.250	1.00 28.40	AAAA
ATOM	1195		YYR Z		44.017	24.215	61.007	1.00 27.52	AAAA
MOTA	1196	CE2	TYR A	A 153	45.216	24.774	61.425	1.00 27.66	AAAA
ATOM	1197	CZ	TYR A	A 153	46.284	24.845	60.547	1.00 28.15	AAAA
	1198	OH	TYR A		47.457	25.407	60.974	1.00 28.83	
ATOM									. AAAA
MOTA	1199	С	TYR A	A 153	42.618	21.172	60.769	1.00 27.82	AAAA
ATOM	1200	0	TYR A	A 153	43.613	20.552	61.110	1.00 27.15	AAAA
ATOM.	1201	N	LEU A	154	41.636	21.487	61.604	1.00 29.25	AAAA.
MOTA	1202	CA	LEU A		41.665	21.138	63.014	1.00 29.35	AAAA
MOTA	1203	CB	LEU A	A 154	40.507	21.829	63.715	1.00 30.25	AAAA
MOTA	1204	CG	LEU A	154	40.685	23.346	63.792	1.00 31.10	AAAA
	1205		LEU A		39.348	24.020	64.092	1.00 31.24	
ATOM					•				AAAA
MOTA	1206		LEU A		41.747	23.669	64.852	1.00 29.84	AAAA
ATOM	1207	C	LEU A	154	41.625	19.639	63.263	1.00 29.73	AAAA
ATOM	1208	0	LEU A	154	42.313	19.151	64.150	1.00 30.51	AAAA
	1209	N	ARG A		40.832	18.903	62.489	1.00 28.95	
MOTA									AAAA
MOTA	1210	CA	ARG A		40.771	17.459	62.671	1.00 28.94	AAAA
ATOM	1211	CB	ARG A	155	39.742	16.820	61.723	1.00 28.64	AAAA
ATOM	1212	CG	ARG A	155	38.312	17.312	61.952	1.00 27.82	AAAA
								1.00 27.19	
ATOM	1213	CD	ARG A		37.319	16.751	60.955		AAAA
ATOM	1214	NE	ARG A	155	36.804	15.444	61.338	1.00 28.86	. AAAA
MOTA	1215	CZ	ARG A	155	35.939	14.742	60.612	1.00 28.93	AAAA
ATOM	1216	NH1	ARG A	155	35.500	15.227	59.459	1.00 29.47	AAAA
	1217		ARG A		35.486	13.574	61.053	1.00 28.76	
ATOM									AAAA
MOTA	1218	С	ARG A	155	42.158	16.853	62.438	1.00 30.20	AAAA
MOTA	1219	0	ARG A	. 155	42.572	15.949	63.164	1.00 30.74	AAAA
ATOM	1220	N	LYS A		42.890	17.362	61.447	1.00 30.32	· AAAA
MOTA	1221	CA	LYS A		44.224	16.838	61.173	1.00 30.07	AAAA
ATOM	1222	CB	LYS A	156	44.771	17.373	59.847	1.00 30.26	AAAA
ATOM .	1223	CG	LYS A	156	46.168	16.869	59.525	1.00 30.16	AAAA
ATOM	1224	CD	LYS A		46.686	17.368	58.181	1.00 31.19	AAAA
ATOM	1225	CE	LYS A		45.884	16.813	56.986	1.00 31.70	AAAA
ATOM	1226	NZ	LYS A	156	45.963	15.324	56.824	1.00 31.20	AAAA
ATOM	1227	С	LYS A	156	45.167	17.202	62.306	1.00 30.08	AAAA
ATOM	1228	ō	LYS A		46.192	16.550	62.485	1.00 29.16	· AAAA
ATOM	1229	N	LYS A		44.816	18.252	63.053	1.00 30.08	AAAA
ATOM	1230	CA	LYS A	157	45.608	18.691	64.196	1.00 31.03	AAAA
ATOM '	1231	CB	LYS A	157	45.446	20.201	64.452	1.00 31.81	AAAA
ATOM	1232	CG	LYS A		46.067	21.134	63.419	1.00 32.12	AAAA
ATOM	1233	CD	LYS A		47.580	21.041	63.348	1.00 31.34	AAAA
MOTA	1234	CE.	LYS A	157	48.080	21.941	62.226	1.00 32.66	AAAA.
MOTA	1235	NZ	LYS A	157	49.556	21.921	61.996	1.00 32.74	\AAA
	1236	С	LYS A		45.196	17.923	65.458	1.00 31.73	. `AAA
MOTA	_								
MOTA	1237	O.	LYS A		45.652	18.230	66.558	1.00 31.93	AAAA
ATOM	1238	N	GLY A	158	44.312	16.942	65.299	1.00 32.41	AAAA
MOTA	1239	CA	GLY A	158	43.901	16.140	66.436	1.00 32.34	AAAA
-			GLY A		42.604	16.429	67.172	1.00 32.65	AAAA
MOTA	1240	C							
MOTA	1241	0	GLY A		42.182	15.604	67.980	1.00 32.85	AAAA
ATOM	1242	N	PHE A	159	41.960	17.565	66.932	1.00 33.16	AAAA
ATOM	1243	CA	PHE A		40.712	17.842	67.650	1.00 34.16	AAAA
MOTA	1244	CB	PHE A		40.220	19.281	67.403	1.00 34.81	AAAA
ATOM	1245	CG	PHE A	159	41.134	20.343	67.965	1.00 34.01	AAAA
ATOM	1246		PHE A		42.327	20.669	67.329	1.00 34.18	AAAA
					40.821		69.166	1.00 34.61	AAAA
MOTA	1247		PHE A			20.981			
MOTA	1248		PHE A		43.197	21.610	67.874	1.00 33.65	AAAA
ATOM '	1249	CE2	PHE A	159	41.689	21.924	69.718	1.00 34.52	AAAA
	1250	CZ	PHE A		42.878	22.236	69.065	1.00 33.90	AAAA
MOTA		25							
ATOM	1251		PHE A		39.645	16.840	67.239	1.00 34.04	AAAA
ATOM	1252	0	PHE A	159	39.568	16.456	66.068	1.00 34.98	AAAA
ATOM	1253	N	LYS A	160	38.839	16.403	68.202	1.00 33.36	AAAA
31011			LYS A		37.794	15.415	67.936	1.00 33.11	AAAA
ATOM	1254	CA	א נינע	200	31.134	-7.473		33.11	ALC: N

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MOTA	1255	CB	LYS	A	160		38.060	14.140	68.763	1.00 33.97	AAAA
MOTA	1256	CG			160		39.410	13.491	68.457	1.00 35.31	AAAA
ATOM	1257	CD	LYS	A	160		39.833	12.364	69.429	1.00 36.48	AAAA
ATOM	1258	CE			160		39.095	11.037	69.243	1.00 37.97	AAAA
MOTA	1259	NZ			160		37.636	11.080	69.568	1.00 39.67	AAAA
MOTA	1260	С			160		36.385	15.941	68.210	1.00 31.68	AAAA
MOTA	1261	0			160		35.405	15.290	67.887	1.00 31.51	AAAA
. ATOM	1262	N			161		36.291	17.114	68.819	1.00 31.11 1.00 30.92	AAAA AAAA
MOTA	1263	CA			161		35.003	17.719 17.592	69.114 70.604	1.00 30.92	AAAA
MOTA	1264	CB			161		34.655 34.451	16.157	71.102	1.00 32.91	AAAA
MOTA	1265 1266	CG CD			161 161		33.994	16.126	72.570	1.00 33.26	AAAA
MOTA	1267	NE			161		34.929	16.797	73.476	1.00 34.01	AAAA
ATOM ATOM	1268	CZ			161		36.183	16.404	73.698	1.00 34.88	AAAA
ATOM	1269		ARG				36.675	15.334	73.081	1.00 34.89	AAAA
ATOM	1270		ARG				36.954	17.084	74.537	1.00 34.71	AAAA
ATOM	1271	C	ARG	Α	161		35.061	19.185	68.714	1.00 30.28	AAAA
MOTA	1272	0	ARG	Α	161		35.365	20.059	69.529	1.00 29.86	AAAA
ATOM	1273	N			162		34.774	19.433	67.437		· AAAA
MOTA	1274	CA			162		34.788	20.774	66.862 65.464	1.00 26.41 1.00 26.87	AAAA AAAA
ATOM	1275	CB			162		35.443 35.453	20.762 22.160	64.872	1.00 26.91	AAAA
MOTA	1276		ILE				36.877	20.234	65.578	1.00 28.19	AAAA
MOTA	1277 1278		ILE				37.614	20.090	64.240	1.00 28.24	AAAA
MOTA MOTA	1279	C			162		33.369	21.283	66.731	1.00 24.08	AAAA
MOTA	1280	ŏ			162		32.485	20.572	66.267	1.00 24.40	AAAA
ATOM	1281	N	LEU	Α	163		33.153	22.519	67.153	1.00 22.25	AAAA
MOTA	1282	CA	LEU				31.838	23.126	67.074	1.00 20.48	AAAA
ATOM	1283	CB	LEU				31.408	23.671	68.440	1.00 20.97	AAAA AAAA
MOTA	1284	CG	LEU				30.099	24.477 23.695	68.486 67.799	1.00 20.50 1.00 19.07	AAAA
MOTA	1285		LEU				28.998 29.738	24.802	69.950	1.00 19.76	AAAA
MOTA	1286	CD2	LEU				31.801	24.241	66.055	1.00 18.76	AAAA
MOTA MOTA	1287 1288	Ö	LEU				32.756	24.986	65.894	1.00 18.41	AAAA
ATOM	1289	N			164		30.677	24.344	65.368	1.00 17.85	AAAA
ATOM	1290	CA	TYR	Α	164		30.496	25.372	64.373	1.00 17.16	AAAA
ATOM	1291	CB	TYR	Α	164		30.644	24.768	62.983	1.00 17.45	AAAA
MOTA	1292	CG	TYR				30.484	25.783	61.900	1.00 17.70	AAAA AAAA
MOTA	1293		TYR					26.772 27.734	61.701 60.721	1.00 16.23 1.00 17.35	AAAA
ATOM .	1294		TYR TYR				31.280 29.350	25.781	61.092	1.00 17.95	AAAA
MOTA	1295 1296		TYR				29.173	26.746	60.103	1.00 18.03	AAAA
MOTA MOTA	1297	CZ	TYR				30.138	27.717	59.919	1.00 17.30	AAAA
ATOM	1298	ОН	TYR				29.955	28.647	58.926	1.00 16.70	AAAA
ATOM	1299	С	TYR				29.123	26.016	64.514	1.00 15.85	AAAA
ATOM	1300	0	TYR	Α	164	•	28.101	25.351	64.416	1.00 16.44	AAAA
ATOM	1301	N	ILE				29.115	27.319	64.743	1.00 15.54	AAAA
MOTA		CA			165	•	27.878	28.088	64.897 66.250	1.00 15.71 1.00 15.18	AAAA AAAA
ATOM	1303	CB	ILE				27.869 26.621	28.819 29.685	66.374	1.00 13.18	AAAA
ATOM	1304		ILE				28.000	27.797	67.386	1.00 13.94	AAAA
ATOM	1305 1306		ILE				28.356	28.421	68.747	1.00 13.94	AAAA
ATOM	1307	C	ILE				27.808	29.124	63.754	1.00 16.00	AAAA
MOTA MOTA	1308	ō	ILE				28.711	29.941	63.576	1.00 16.56	AAAA
ATOM	1309	N	ASP			*	26.721	29.087	63.001	1.00 16.18	AAAA
ATOM	1310	CA	ASP				26.524	29.962	61.865	1.00 16.67	AAAA
ATOM	1311	CB	ASP				26.240	29.066	60.651	1.00 18.05	AAAA
ATOM	1312	CG	ASP				26.238	29.809	59.329	1.00 19.21	AAAA AAAA
MOTA	1313		ASP				25.353	30.659	59.114 58.495	1.00 18.36 1.00 19.19	AAAA
ATOM	1314		ASP				27.131	29.521	62.169	1.00 17.57	AAAA
ATOM	1315	C	ASP				25.342 24.206	30.904 30.459	62.321	1.00 17.26	AAAA
ATOM	1316	O N	ASP LEU				25.605	32.202	62.274	1.00 16.67	AAAA
ATOM	1317	CA	LEU				24.526	33.135	62.562	1.00 16.89	AAAA
MOTA	1318 1319	CB	LEU				24.923	34.116	63.663	1.00 17.27	AAAA
ATOM ATOM	1320	CG	LEU				25.499	33.529	64.954	1.00 18.37	AAAA
ATOM				-					•		

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ATOM	1321	CD1 LEU A 167	25.760	34.671	65.933	1.00 18.72	AAAA
MOTA	1322	CD2 LEU A 167	24.566				AAAA
MOTA	1323	C LEU A 167	24.146				AAAA
MOTA	1324	O LEU A 167	23.390			1.00 17.21	AAAA
ATOM	1325	N ASP A 168	24.683			1.00 17.83	AAAA
ATOM	1326	CA ASP A 168	24.382	-		1.00 17.84	AAAA
MOTA	1327	CB ASP A 168	25.178			1.00 20.42	AA-AA
MOTA	1328	CG ASP A 168	25,140			1.00 21.41	AAAA
ATOM	1329	C ASP A. 168	22.915		58.660	1.00 18.35	AAAA
MOTA	1330	O ASP A 168	22.419			1.00 19.62	AAAA
MOTA	1331	OD1 ASP A 168	- 26.066		56.330	1.00 22.42	AAAA
ATOM		OD2 ASP A 168	24.186		55.746	1.00 21.79	AAAA
MOTA.	1333	N ALA A 169	22.239		58.010	1.00 17.98	AAAA
MOTA	1334	CA ALA A 169	20.824			1.00 17.36	AAAA
MOTA	1335	CB ALA A 169	20.348		57.007	1.00 17.00	AAAA
MOTA	1336	C ALA A 169	20.439		56.887	1.00 18.64	AAAA
MOTA	1337	O ALA A 169	19.255	33.043	56.819	1.00 19.46	AAAA
MOTA	1338	N HIS A 170	21.412	32.712	56.262	1.00 18.71	AAAA
MOTA	1339	CA HIS A 170	21.107	31.518	55.464	1.00 18.43	AAAA
MOTA	1340	C HİS A 170	21.802	30.265	55.986	1.00 18.02	AAAA
ATOM	1341	O HIS A 170	22.910	30.332	56.514	1.00 17.20	AAAA
ATOM	1342	CB HIS A 170	21.539	31.678	54.004	1.00 18.79	AAAA
ATOM		CG HIS A 170	21.137	32.968	53.386	1.00 17.65	AAAA
ATOM	1344 1	ND1 HIS A.170	21.644	34.162	53.828	1.00 18.08	AAAA
ATOM		CE1 HIS A 170	21.112	35.081	53.054	1.00 18.95	AAAA
ATOM		CD2 HIS A 170	20.301	33.194	52.348	1.00 18.81	AAAA
MOTA		VE2 HIS A 170	20.291	34.544	52.140	1.00 19.66	AAAA
MOTA		N HIS A 171	21.142	29.124	55.793	1.00 17.53	AAAA
ATOM	-	CA HIS A 171	21.662	27.822	56.193	1.00 16.38	AAAA
ATOM	-	CB HIS A 171	20.644	26.740	55.830	1.00 16.32	AAAA
ATOM		G HIS A 171	21.157	25.337	55.958	1.00 15.91	AAAA
MOTA		D2 HIS A 171	21.241	24.336	55.051	1.00 14.14	AAAA
ATOM		TD1 HIS A 171	21.602	24.807	57.151	1.00 16.79	AAAA
MOTA		E1 HIS A 171	21.937	23.543	56.973	1.00 14.91	AAAA
MOTA		TE2 HIS A 171	21.725	23.234	55.709	1.00 15.45	AAAA
MOTA	1356 C		22.982	27.522	55.509	1.00 16.94	AAAA
ATOM ATOM	1357 C		23.146 23.926	27.725	54.318	1.00 18.71	AAAA
MOTA		A CYS A 172	25.237	27.019	56.279 55.778	1.00 16.99	AAAA.
ATOM		B CYS A 172	26.219	26.670 26.721	56.947	1.00 16.23 1.00 17.89	AAAA
ATOM		G CYS A 172	25.638	25.773	58.397	1.00 17.89	AAAA
ATOM	1362 C		25.205	25.271	55.210	1.00 17.89	AAAA AAAA
ATOM	1363 0		25.947	24.413	55.670	1.00 17.66	AAAA
MOTA	1364 N		24.364	25.026	54.214	1.00 18.25	AAAA
ATOM	1365 C		24.253	23.680	53.620	1.00 19.91	AAAA
ATOM	1366 C		23.342	23.699	52.397	1.00 20.86	AAAA
ATOM	1367 C	G ASP A 173	23.780		51.358	1.00 21.90	AAAA
ATOM	1368 0	D1 ASP A 173	23.257	24.640	50.217	1.00 21.35	AAAA
ATOM	. 1369 0	D2 ASP A 173	24.624	25.597	51.687	1.00 21.35	AAAA
ATOM	1370 C	ASP A 173	25.573	23.021	53.227	1.00 21.02	AAAA
ATOM	1371 0	ASP A 173	25.673	21.785	53.199	1.00 22.79	AAAA
ATOM	1372 N	GLY A 174	26.579	23.832	52.912	1.00 20.03	AAAA
ATOM	1373 C		27.870	23.277	52.553	1.00 19.72	AAAA
ATOM	1374 C	GLY A 174	28.537	22.680	53.771	1.00 20.27	AAAA
MOTA	1375 0	GLY A 174	29.110	21.599	53.711	1.00 19.77	AAAA
ATOM	1376 N	VAL A 175	28.448	23.387	54.893	1.00 21.38	AAAA
MOTA	1377 C		29.056	22.934	56.135	1.00 22.26	AAAA
MOTA	1378 CE		29.032	24.040	57.203	1.00 23.15	AAAA
ATOM		31 VAL A 175	29.853	23.617	58.418	1.00 22.84	AAAA
ATCM		32 VAL A 175	29.562		.56.612	1.00 23.43	AAAA
MOTA	1381 C	VAL A 175	28.302	21.724	56.654	1.00 23.51	AAAA
ATOM	1382 0	VAL A 175	28.893	20.803	57.210	1.00 23.74	AAAA
ATOM	1383 N	GLN A 176	26.993	21.721	56.452	1.00 24.80	AAAA
MOTA	1384 CA		26.171	20.601	56.893	1.00 25.41	AAAA
MOTA	1385 CE		24.689	20.913	56.694	1.00 24.77	AAAA
ATOM .	1386 CG	GLN A 176	23.799	19.735	57.036	1.00 26.23	AAAA
							_

								•	
ATOM	1387	CD	GLN A	176	22.33	34 20.094	57.069	1.00 27.17	AAAA
	1388		GLN A		21.90		57.911	1.00 28.24	AAAA
ATOM		NE2			21.5		56.151	1.00 26.54	AAAA
MOTA	1389							•	
ATOM	1390	С	GLN A		26.5		56.180	1.00 25.67	AAAA
ATOM	1391	0	GLN A	176	26.7		56.820	1.00 26.98	AAAA
MOTA	1392	N	GLU A	177	26.49		54.853	1.00 26.34	AAAA
MOTA	1393	CA	GLU A	177	26.78		54.073	1.00 26.18	AAAA
MOTA	1394	CB	GLU A		26.7		52.580	1.00 .27.13	AAAA
			GLU A		26.7		51.628	1.00 29.77	AAAA
ATOM	1395	CG							
MOTA	1396		_GLU A		26.63		50.147	1.00 31.29	AAAA
ATOM	1397		GLU A		27.6		49.500	1.00 31.35	АДДА
MOTA	1398	OE2	GLU A	177	25.4	71 17.703	49.636	1.00 32.05	AAAA
ATOM	1399	С	GLU A	177	28.1	60 17.556	54.460	1.00 26.38	AAAA
ATOM	1400	0	GLU A	177	28.33	38 16.349	54.595	1.00 25.41	AAAA
ATOM	1401	N	ALA A		29.1		54.659	1.00 -27.58	AAAA
	1402	CA	ALA A		30.49		55.004	1.00 27.66	- AAAA
MOTA							54.994	1.00 26.20	
MOTA	1403	CB	ALA A		. 31.3				AAAA
MOTA	1404	C	ALA A				56.318	1.00 28.33	AAAA
MOTA	1405	0	ALA A	178	31.6		56.439	1.00 29.44	AAAA
ATOM	1406	N	PHE A	179	29.8	49 17.564	57.308	1.00 28.25	AAAA
ATOM	1407	CA	PHE A	179	30.03	36 16.852	58.561	1.00 29.20	AAAA
MOTA	1408	CB	PHE A	179	30.5	70 17.794	59.624	1.00 29.35	AAAA
	1409	CG	PHE A		31.7		59.171	1.00 30.26	AAAA
MOTA			PHE A		31.5		58.497	1.00 31.01	AAAA
MOTA	1410							1.00 30.37	
MOTA	1411		PHE A		33.0		59.339		AAAA
MOTA	1412		PHE A		32.6		57.993	1.00 31.20	AAAA
MOTA	1413	CE2	PHE A	179	34.13		58.840	1.00 31.74	AAAA
ATOM	1414	CZ	PHE A	179	33.9	50 19.960	58.161	1.00 31.81	AAAA
ATOM	1415	С	PHE A	179	28.7	60 16.180	59.040	1.00 30.33	AAAA
ATOM	1416	0	PHE A	179	28.6	24 15.810	60.215	1.00 31.82	AAAA
MOTA	1417	N	TYR A				58.105	1.00 29.18	AAAA
	1418	CA	TYR A		26.5		58.389	1.00 28.99	AAAA
ATOM					25.7		57.123	1.00 28.30	AAAA
MOTA	1419	CB	TYR A					1.00 28.27	
ATOM	1420	CG	TYR A				57.384		AAAA
MOTA	1421		TYR A		23.3		57.021	1.00 27.05	AAAA
ATOM	1422	CEI	TYR A	180	22.0		57.197	1.00 28.51	AAAA
ATOM	1423	CD2	TYR A	180	23.6		57.942	1.00 29.34	AAAA
MOTA	1424	CE2	TYR A	180	22.3	16 16.691	58.125	1.00 29.27	AAAA
ATOM	1425	CZ	TYR A	180	21.4	95 15.645	57.746	1.00 29.33	AAAA
ATOM	1426	OH	TYR A		20.1	41 15.775	57.893	1.00 30.83	AAAA
ATOM	1427	C	TYR A				58.940	1.00 28.79	AAAA
	1428	õ	TYR A		25.8		59.785	1.00 28.71	AAAA
MOTA		N	ASP A		27.6		58.472	1.00 29.03	AAAA
MOTA	1429						58.914	1.00 28.49	AAAA
MOTA	1430	CA	ASP A		27.8			1.00 27.59	AAAA
ATOM	1431	CB	ASP A		28.1				
MOTA	1432	CG	ASP A		29.5		57.229	1.00 28.82	AAAA
ATOM	1433	OD1	ASP A	181	29:9		57.183	1.00 29.25	AAAA
ATOM	1434	OD2	ASP A	181	30.2	16 .10 119	56.887	1.00 28.68	AAAA
ATOM	1435	C	ASP A	181	28.8	63 11.631	60.009	1.00 27.67	AAAA
ATOM	1436	0	ASP A	181	29.2	71 10.504	60.293	1.00 27.57	AAAA
	1437	N	THR A			08 12.713		1.00 26.27	AAAA
MOTA	1430		THR A		30.2	• •	61.689	1.00 26.22	AAAA
HOTA	1438	CA					61.317	1.00 25.92	AAAA
ATOM	1439	CB	THR A		31.6				
ATOM	1440		THR A		32.5		62.416	1.00 25.06	AAAA
MOTA	1441	CG2	THR A		31.5		60.974	1.00 25.25	AAAA
ATCM	1442	С	THR A		29.7		62.934	1.00 25.87	AAAA
ATOM	1443	0	THR A	182	28.9	42 14.102	62.863	1.00 26.35	AAAA
MOTA	1444	N	ASP A		30.3		64.071	1.00 25.86	AAAA
	1445	CA	ASP A		29.9		65.355	1.00 26.12	AAAA
ATOM			ASP A		29.4		66.274	1.00 27.10	AAAA
ATOM	1446	CB					66.488	1.00 28.80	AAAA
atch	1447	CG	ASP A		30.5				AAAA
ATOM	1448	OD1	ASP A	193	31.0		65.480	1.00 29.78	
ATOM	1449		ASP A		30.7		67.657	1.00 29.21	AAAA
ATOM	1450	С	ASP A		31.1			1.00 26.77	AAAA
ATOM	1451	0	ASP A	183	31.0		67.146	1.00 26.65	AAAA
TOM	1452	N	GLN A		32.2		65.254	1.00 26.05	AAAA

1452 N GLN A 184 32.229 14.254 65.254 1.00 26.05

SUBSTITUTE SHEET (RULE 26)

3	1453	-					_		
MOTA	1453		GLN A		33.381	14.983	65.766	1.00 25.	.85 AAAA
ATOM	1454	CB	GLN A	184	34.674	14.510	65.095	1.00 26.	AAAA 08
ATOM	1455	CG	GLN A	184	34.920		65.303	1.00 27.	
ATOM	1456	CD	GLN A		36.273				
ATOM	1457		GLN A				64.822	1.00 28.	40 адад
					36.685		63.709	1.00 30.	
MOTA	1458	NE2	GLN A		36.970	11.816	65.651	1.00 29.	24 AAAA
MOTA	1459	С	GLN A	184	33.159	16.474	65.536	1.00 25.	
ATOM	1460	0	GLN A	184	33.734		66.220	1.00 24	
ATOM	1461	N	VAL A						
					32.290	16.791	64.584	1.00 25.	
ATOM	1462	CA	VAL A		31.975	18.182	64.291	1.00 24.	
ATOM '	1463	CB	VAL A	185	32.324	18.563	62.832	1.00 23.	20 AAAA
MOTA	1464	CG1	VAL A	185	32.045	20.060	62.599	1.00 19.	
ATOM	1465		VAL A		33.777	18.205	62.543		
ATOM	1466	c	VAL A					1.00 20.	
					30.494	18.421	64.501	1.00 24.	
MOTA	1467	0	VAL A		29.664	17.787	63.844	1.00 27.	07 дада
MOTA	1468	N	PHE A	186	30.162	19.311	65.434	1.00 23.	
ATOM	1469	CA	PHE A	186	28.768	19.645	65.684	1.00 20.	
ATOM	1470	CB	PHE A	186	28.513	19.937	67.164	1.00 19.	_
ATOM	1471	CG	PHE A		27.057	20.037	67.500		
	1472							1.00 18.	
ATOM			PHE A		26.359	18.918		1.00 17.	
MOTA	1473		PHE A		26.358	21.213	67.263	1.00 17.	46 AAAA
MOTA	1474		PHE A		24.999	18.964	68.147	1.00 17.	
MOTA	1475	CE2	PHE A	186	24.997	21.271	67.459	1.00 18.	
MOTA	1476	CZ	PHE A			.20.138	67.905	1.00 18.	
ATOM	1477	c	PHE A		28.464	20.911			
	1478	ō					64.895	1.00 19.	
MOTA			PHE A		29.079	21.940	65.129	1.00 18.	
ATOM	1479	N	VAL A		27.520	20.834	63.964	1.00 18.	34 AAAA
ATOM	1480	CA	VAL A		27.137	21.993	63.160	1.00 16.	47 AAAA
ATOM	1481	CB	VAL A	187	27.006	21.630	61.655	1.00 14.	
ATOM	1482	CG1	VAL A	187	26.628	22.869	60.828	1.00 10.	
ATOM	1483	CG2	VAL A	187	28.314	21.031	61.160	1.00 12.	
MOTA	1484	С	VAL A		25.806	22.511	63.665		
ATOM	1485	ŏ	VAL A					1.00 17.	
					24.852	21.746	63.792	1.00 16.	
ATOM.	1486		LEU A		25.763	23.809	63.960	1.00 18.	
MOTA	1487	CA	LEU A		24.555	24.507	64.460	1.00 20.	51 AAAA
MOTA	1488	CB	LEU A	188	24.752	24.995	65.914	1.00 21.	24 AAAA
ATOM	1489	CG	LEU A 1	188	23.702	26.019	66.395	1.00 20.	
MOTA	1490	CD1	LEU A 1	188	22.365	25.323	66.493	1.00 19.	
ATOM	1491	CD2	LEU A 1	188	24.085	26.627	67.750	1.00 20.	
ATOM	1492	С	LEU A		24.297	25.735	63.591		
ATOM	1493	ŏ	LEU A					1.00 20.	41 AAAA
					25.223	26.484	63.288	1.00 21.	
MOTA	1494	N	SER A 1		23.049	25.987	63.233	1.00 19.	
MOTA	1495	CA	SER A 1		22.786	27.130	62.381	1.00 18.	aaaa do
ATOM	1496	CB	SER A I	189	22.970	26.715	60.906	1.00 18.	54 AAAA
ATOM	1497	OG	SER A 1	189	22.559	27.731	59.998	1.00 17.	
MO7 A	1498	С	SER A 1	89	21.418	27.751	62.554	1.00 17.	
MO'LE	1499	ō	SER A 1		20.404	27.051	62.540		
								1.00 19.	
A_OM	1500	N	LEU A 1		21.386	29.067	62.722	1.00 16.	
ATOM	1501		LEU A 1		20.117	29.772	62.797	1.00 18.	49 AAAA
ATOM	1502	CB	LEU A 1	.90	20.097	30.865	63.886	1.00 17.	
ATOM	1503	CG	LEU A 1	.90	20.534	30.600	65.337	1.00 17.	·
ATOM	1504	CD1	LEU A 1		19.643	31.406		1.00 15.	
ATOM	1505		LEU A 1		20.455	29.147		1 00 15.	50 AAAA
							65.686	1.00 15.	
MOTA	1506		LEU A 1		20.111	30.408	61.416	1.00 19.	
MOTA	1507		LEU A 1		21.136	30.891	60.967	1.00 19.	75 AAAA
atom	1508	N	HIS A 1	91	18.975	30.397	60.736	1.00 21.	75 AAAA
MOTA	1509	ÇA	HIS A 1	91	18.897	30.955	59.383	1.00 23.	
ATOM	1510	CB	HIS A 1	91	19.626	30.013	58.426	1.00 23.0	
ATOM	1511		HIS A 1		19.157	28.597	58.533		
	1512		HIS A 1		19.770	27.485		1.00 24.2	
MOTA							59.009	1.00 23.	
ATOM	1513		HIS A 1		17.869	28.217	58.217	1.00 24.7	
ATOM	1514		HIS A 1		17.709	26,935	58.491	1.00 23.9	аааа ое
ATOM	1515		HIS A 1		18.849	26.467	58.973	1.00 24.5	
MOTA	1516	C :	HIS A 1	91	17.446	31.119	58.926	1.00 24.	
ATOM	1517		HIS A 1		16.519	30.658	59.596	1.00 24.9	
ATOM	1518		GLN A 1		17.249	31.789	57.794	1.00 24.3	
TTOM.	1210	•• '			21.447	31.707	- 134	1.00 24.3	AAAA EE

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### Figure 19-24

MOTA	1519	CA	GLN A	192	15.899	31.959	57.269	1.00 25.77	AAAA
ATOM	1520	CB	GLN A		15.881	32.896			
			_				56.060	1.00 26.51	
ATOM	1521	CG	GLN A	192	16.467	34.271	56.325	1.00 26.99	AAAA
ATOM	1522	CD	GLN A	192	16.581	35.076	55.062	1.00 27.98	
ATOM	1523	OE1	GLN A	192	15.583	35.496	54.493		
								1.00 30.48	
MOTA	1524	NES	GLN A	192	17.802	35.274	54.595	1.00 29.04	AAAA
ATOM	1525	С	GLN A	192	15.463	30.573	56.832	1.00 25.77	
ATOM	1526	ō	GLN A						
					16.211	29.865	56.169	1.00 26.73	
MOTA	1527	N	SER A	193	14.259	30.184	57.214	1.00 25.48	AAAA
ATOM	1528	CA	SER A	193	13.750	28.877	56.863	1.00 24.51	
ATOM	1529	CB	SER A		12.288	28.788	57.286	1.00 23.77	AAAA
ATOM	1530	OG	SER A	193	11.753	27.517	57.010	1.00 24.81	AAAA
MOTA	1531	С	SER A	193	13.906	28.597	55.361	1.00 24.53	
	1532							•	
ATOM		0	SER A		13.736	29.479	54.522	1.00 22.32	AAAA
MOTA	1533	N	PRO A	194	14.226	27.348	55.007	1.00 25.69	AAAA
ATOM	1534	CD	PRO A	194	14.411	26.167	55.862	1.00 25.02	
ATOM	1535	CA	PRO A				_		
					14.399	26.976	53.604	1.00 27.05	
MOTA	1536	CB	PRO A	194	14.906	25.535	53.697	1.00 26.30	AAAA
MOTA	1537	CG	PRO A	194	15.479	25.466	55.124	1.00 26.44	
MOTA	1538	C	PRO A		13.076	27.057			
							52.849	1.00 27.79	
MOTA	1539	0	PRO A	194	13.066	27.057	51.625	1.00 28.82	AAAA
ATOM	1540	N	GLU A	195	11.966	27.133	53.582	1.00 28.29	AAAA
MOTA	1541	CA	GLU A		10.656	27.187	52.950	1.00 29.08	
MOTA	1542	CB	GLU A		9.534	27.030	54.001	1.00 31.08	AAAA
MOTA	1543	CG	GLU A	195	9.070	28.294	54.722	1.00 35.07	AAAA
ATOM	1544	CD	GLU A	195	7.850	28.980	54.064	1.00 38.05	
ATOM	1545		GLU A		7.389	30.017	54.601	1.00 38.80	
MOTA	1546	OE2	GLU A	195	7.342	28.487	53.024	1.00 39.20	AAAA
ATOM	1547	С	GLU A	195	10.483	28.471	52.150	1.00 28.05	
	1548	ŏ							
MOTA			GLU A		9.722	28.512	51.189	1.00 28.57	
ATOM	1549	N	TYR A	196	11.223	29.510	52.514	1.00 27.39	AAAA
ATOM	1550	CA	TYR A	196	11.108	30.769	51.802	1.00 25.80	
ATOM	1551	CB	TYR A		10.275	31.743	52.645	1.00 24.97	
MOTA	1552	CG	TYR A		10.971	32.281	53.868	1.00 23.41	AAAA
ATOM	1553	CD1	TYR A	196	11.911	33.306	53.765	1.00 23.99	AAAA
ATOM	1554	CEI	TYR A	196	12.559	33.805	54.892	1.00 23.44	
MOTA	1555		TYR A		10.697	31.768	55.126	1.00 23.24	
ATOM	1556	CE2	TYR A	196	11.336	32.256	56.254	1.00 23.93	AAAA
ATOM	1557	CZ	TYR A	196	12.265	33.270	56.133	1.00 24.07	
ATOM	1558	OH	TYR A		12.913	33.731	57.247		
								1.00 25.06	
ATOM	1559	С	TYR A	196	12.450	31.406	51.411	1.00 24.97	AAAA
ATOM	1560	0	TYR A	196	12.475	32.495	50.840	1.00 25.14	AAAA
ATOM	1561	N	ALA A	197	13.563	30.737	51.686	1.00 23.81	
MOTA	1562	CA	ALA A		14.855	31.330	51.337	1.00 23.32	
ATOM	1563	CB	ALA A	197	15.350	32.220	52.488	1.00 23.33	AAAA
ATOM -	1564	С	ALA A	197	15.952	30.356	50.957	1.00 22.74	
	1565		ALA A		15.951		.51 .37 '		
ATOM								1.00 22.47	
- MOTA	1566	N	.PHE A	198	16, 900	30.852	50.16	1.00 23.23	
ATOM	1567	CA	PHE A	198	18.062	30.081	49.741	1.00 23.68	AAAA
ATOM	1568	CB	PHE A		19.083	31.006	49.069	1.00 23.33	
								_	AAAA
MOTA	1569	CG					48.464		AAAA
ATOM	1570	CD1	PHE A	198	20.151	29.713	47.203	1.00 22.75	AAAA
ATOM	1571		PHE A		21.436	30.127	49.175	1.00 23.32	
									AAAA
ATOM	1572		PHE A		21.207	29.003	46.645	1.00 22.13	AAAA
MOTA	1573	CE2	PHE A	198	22.512	29.408	48.622	1.00 22.83	AAAA
ATOM	1574	CZ	PHE A		22.386	28.849	47.351	1.00 22.55	· AAAA
ATOM	1575	С	PHE A		18.689	29.490	51.008	1.00 23.69	AAAA
ATOM	1576	0	PHE A	198	18.802	30.171	52.012	1.00 22.85	AAAA
MOTA	1577	N	PRO A		19.166	28.236	50.954	1.00 23.96	AAAA
MOTA	1578	CD	PRO A		19.833	27.639	52.123	1.00 24.26	AAAA
ATOM	1579	CA	PRO A	199	19.199	27.286	49.837	1.00 24.70	AAAA
ATOM	1580	CB	PRO A		20.163	26.222	50.357	1.00 23.30	AAAA
								1.00 23.21	
ATOM	1581	CG	PRO A		19.797	26.162	51.782	1.00 23.21	AAAA
ATOM	1582	С	PRO A		17.885	26.679	49.326	1.00 25.22	AAAA
ATOM	1583	0	PRO A	199	17.866	26.145	48.215	1.00 26.24	AAAA
	1584	N	PHE A		16.811	26.756	50.116	1.00 25.09	
ATOM	エンロボ	**			10.011	40.750		23.09	AAAA

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					•					
MOTA	1585	CA	PHE A	200		15.497	26.190	49.763	1.00 26.2 <del>9</del>	AAAA
			PHE A			15.064		48.340	1.00 25.65	AAAA
ATOM	1586	CB	PDE A	200				48.122	1.00 24.65	AAAA
MOTA	1587	CG	PHE A			14.863	28.035			
ATOM	1588	CD1	PHE A	200		15.806	28.781	47.439	1.00 24.42	AAAA
MOTA	1589	CD2	PHE A	200		13.735	28.671	48.608	1.00 23.79	AAAA
			PHE A			15.631	30.125	47.246	1.00 24.41	AAAA
ATOM_	1590									AAAA
MOTA	1591	CE2	PHE A	200		13.552	30.035	48.418	1.00 24.94	
MOTA	1592	CZ	PHE A	200		14.499	30.760	47.738	1.00 24.57	'AAAA'
	1593	C	PHE A			15.415	24.656	49.863	1.00 28.54	AAAA
MOTA			PHE A			14.386	24.096	50.251	1.00 28.76	AAAA
MOTA	1594	0								AAAA
MOTA	1595	N	GLU A	201		16.499	23.981	49.504	1.00 29.67	
ATOM .	1596	CA	GLU A	201		16.539	22.528	49.524	1.00 31.88	AAAA
ATOM	1597	CB	GLU A	201	-	17.434	22.045	48.392-	1.00 32.71	AAAA
			GLU A			16.897		47.017	1.00 34.87	AAAA
MOTA	1598	CG				17.898			1.00 35.14	AAAA
MOTA	1599	CD	GLU A	201		17.898	22.147			
MOTA	1600	OE1	GLU A GLU A	201		18.299	20.982	45.735	1.00 36.09	- AAAA
ATOM	1601	OE2	GLU A	201		18.286	· 23.112	45.221	1.00 36.30	AAAA
		c	GLU A	201		16.997	21.894	50.835	1.00 32.77	AAAA
MOTA	1602						20.690		1.00 33.44	AAAA
ATOM	1603	0	GLU A			16.806				
MOTA	1604	N	LYS A	202		17.599	22.690	51.711	1.00 32.31	AAAA
ATOM	1605	CA	LYS A	202		18.101	22.168	52.974	1.00 32.09	AAAA
	1606	СВ	LYS A			19.565	21.750	52.811	1.00 33.02	AAAA
MOTA						19.836	20.847		1.00.34.95	AAAA
MOTA	1607	CG	LYS A							
ATOM	1608	CD	LYS A	202		21.334	20.619		1.00 37.92	AAAA
ATOM	1609	CE	LYS A	202		21.655	19.804	50.169	1.00 39.19	AAAA
ATOM	1610	NZ	LYS A			23.120	19.522	49.988	1.00 38.58	AAAA
			LYS A			17.995		54.037	1.00 30.85	AAAA
MOTA	1611	C						53.739	1.00 30.49	AAAA
MOTA	1612	0	LYS A		٠	17.706	24.389			
MOTA	1613	N	GLY A	203		18.238	22.867		1.00 30.81	AAAA
ATOM	1614	CA	GLY A	203		18.159	23.831	56.356	1.00 30.86	AAAA
	1615	C	GLY A			16.991	23.578	57.280	1.00 30.84	AAAA
MOTA						16.828	24.285		1.00 31.58	AAAA
MOTA	1616	0	GLY A						1.00 30.54	AAAA
MOTA	1617	И	PHE A	204		16.182	22.570			
ATOM	1618	CA	PHE A	204		15.025	22.241		1.00 30.51	AAAA
MOTA	1619	CB	PHE A	204		14.061	21.317	57.058	1.00 29.06	AAAA
	1620	CG	PHE A			13.524	21.890	55.787	1.00 27.13	AAAA
MOTA						14.222	21.762	54.601	1.00 26.52	AAAA
MOTA	1621		PHE A						1.00 26.50	AAAA
ATOM	1622		PHE A			12.307	22.548			
ATOM	1623	CE1	PHE A	204		13.713	22.276		1.00 26.44	AAAA
ATOM	1624	CE2	PHE A	204		11.786	23.069	54.600	1.00 26.69	AAAA
	1625	CZ	PHE A			12.490	22.931	53.416	1.00 25.65	AAAA
ATOM						15.401	21.590	59.127	1.00 30.87	AAAA
atom	1626	С	PHE A							AAAA
MOTA	1627	0	PHE A	204		16.395	20.875	59.228	1.00 31.12	
ATOM	1628	N	LEU A	205		14.580	21.844		1.00 31.22	AAAA
ATOM	1629	CA	LEU A	205		14.782	21.329	61.489	1.00 31.43	AAAA
			LEU A			13.575	21.691	62.357	1.00 31.42	AAAA
ATOM	1630					13.603	21.078		1.00 31.76	AAAA:
ATOM	1631	CG	LEU A			_			_	AAAA
ATOM	1632		LEU A			14.894	21.492		1.00 32.36	
ATOM	1633	CD2	LEU A	205		12.379	21.516	64.536	1.00 31.31	AAAA
	1634	С	LEU A			15.026	19.829	61.625	1.00 31.35	AAAA
ATOM						15.714	19.392		1.00 31.40	AAAA
MOTA	1635	0					10.050	60.707	1.00 31.79	AAAA
ATOM	1636	N	GLU A	206		14.448	19.059			
ATOM	1637	CA	GLU A	206		14.509	17.603	60.706	1.00 32.08	AAAA
ATOM	1638	CB	GLU A	206		13,485	17.054	59.716	1.00 33.18	AAAA
		ĊG	GLU A			12,069	17.651	59.829	1.00 34.20	AAAA
MOŢA	1639						19.136	59.453	1.00 33.44	AAAA
ATOM	1640	CD	GLU A			11.973				AAAA
ATOM	1641	OE1	GLU A	206		10.854	19.675	59.422	1.00 33.32	
ATOM	1642	OE2	GLU A	206		13.005	19.777	59.194	1.00 35.12	AAAA
	1643	c	GLU A			15.882	17.045	60.363	1.00 32.34	AAAA
ATOM						16.209	15.909	60.706	1.00 31.83	AAAA
MOTA	1644	0	GLU A							AAAA
MOTA	1645	N	GLU A	207		16.680	17.847	59.670	1.00 32.48	
ATOM	1646	CA	GLU A	207		18.017	17.431	59.287	1.00 31.67	AAAA
	1647	CB	GLU A			18.552	18.385	58.238	1.00 30.39	AAAA
MOTA						17.768	18.316	56.960	1.00 29.63	AAAA
MOTA	1648	CG	GLU A						1.00 30.04	AAAA
ATOM	1649	CD	GLU A			17.953	19.547	56.121	1.00 30.04	
ATOM	1650	OE1	GLU A	207		19.108	19.991	55.971	1.00 30.31	AAAA
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ATOM	1651	OE2	GLU	А	207	16.947	20.070	55.604	1.00 30.76	AAAA
ATOM	1652	С			207	18.879	17.433	60.537	1.00 32.04	AAAA
ATOM	1653	ō			207	19.472	18.448	60.910	1.00 31.57	AAAA
MOTA	1654	N			208	18.935	16.272	61.178	1.00 32.57	AAAA
	1655	CA			208	19.674	16.111	62.408	1.00 33.37	AAAA
ATOM					208			63.519	1.00 33.65	
ATOM	1656	CB				18.709	15.647			AAAA
ATOM	1657		ILE			19.443	15.380	64.806	1.00 34.11	AAAA
MOTA	1658		ILE			17.673	16.742	63.757	1.00 33.94	AAAA
ATOM	1659		ILE			16.628	16.386	64.794	1.00 37.00	AAAA
MOTA	1660	C			208	20.863	15.174	62.280	1.00 34.00	AAAA
ATOM	1661	0			208	21.506	14.829	63.265	1.00 34.40	AAAA
MOTA	1662	N	GLY	Α	209	21.177	14.768	61.062	1.00 34.64	AAAA
ATOM .	1663	CA	GLY	Α	209	22.321	13.903	60.913	1.00 35.55	AAAA
MOTA	1664	С	GLY	À	209	22.164	12.671	60.057	1.00 36.80	AAAA
ATOM	1665	0	GLY	Α	209	21.148	12.461	59.400	1.00 37.32	AAAA
ATOM	1666	N	GLU	A	210	23.199	11.836	60.100	1.00 37.78	AAAA
ATOM	1667	CA			210	23.256	10.621	59.315	1.00 38.04	AAAA
ATOM	1668	CB			210	23.600	11.013	57.892	1.00 38.54	AAAA
ATOM	1669	CG			210	23.469	9.960	56.858	1.00 38.99	AAAA
ATOM	1670	CD			210	24.118	10.412	55.580	1.00 40.10	AAAA
MOTA	1671		GLU			25.365	10.437	55.555	1.00 40.86	AAAA
ATOM	1672		GLU			23.396	10.767	54.619	1.00 40.41	AAAA
	1673	C			210	24.377	9.770	59.894	1.00 37.98	AAAA
MOTA	1674	Ö			210	25.498	10.244	60.041	1.00 38.52	AAAA
ATOM		Ŋ			211	24.085	8.517	60.220	1.00 38.02	AAAA
MOTA	1675					25.116	7.654	60.770	1.00 38.09	AAAA
ATOM	1676	CA			211		8.075	62.164	1.00 38.26	AAAA
ATOM	1677	C			211 211	25.542 24.697	8.443	62.104	1.00 38.20	AAAA
MOTA	1678	0						62.434	1.00 37.82	
MOTA	1679	N			212	26.848	8.030	63.743	1.00 38.20	AAAA
MOTA	1680	CA	LYS			27.396	8.399		1.00 37.36	AAAA
ATOM	1681	CB			212	28.921	8.209	63.766		AAAA
ATOM	1682	CG			212	29.416	6.810	63.385	1.00 40.93	AAAA
ATOM	1683	CD			212	29.001	5.746	64.405	1.00 42.04	AAAA
MOTA	1684	CE			212	29.251	4.318	63.891	1.00 42.80	AAAA
	1685	NZ			212	30.673	4.002	63.562	1.00 42.32	AAAA
MOTA	1686	С			212	27.093	9.859	64.054	1.00 37.08	AAAA
ATOM	1687	0			212	27.075	10.269	65.218	1.00 36.94	AAAA
MOTA	1688	N			213	26.854	10.636	63.002	1.00 35.41	AAAA
MOTA	1689	CA			213	26.592	12.054	63.170	1.00 34.24	AAAA
MOTA	1690	C			213	25.163	12.438	63.470	1.00 33.27	AAAA
ATOM	1691	0			213	24.861	13.611	63.666	1.00 33.29	AAAA
ATOM	1692	N			214	24.280	11.451	63.512	1.00 31.79	AAAA
ATOM	1693	CA			214	22.883	11.710	63.794	1.00 30.47	AAAA
MOTA	1694	CB			214	22.111	10.396	63.737	1.00 30.35	AAAA
MOTA	1695	CG			214	20.676	10.552	63.280	1.00 30.45	AAAA
MOTA	1696	CD			214	20.141	9.241	62.759	1.00 29.75	AAAA
ATOM	1697	CE	LYS	A	214	18.737	9.400	62.229	1.00 30.19	AAAA
ATOM	1698	NZ			214	18.179	8.138	61.671	1.00 31.35	AAAA
ATOM	1699	C	LYS	A	214	22.778	12.374	65.168	1.00 30.31	AAAA
ATOM	1700	0	LYS	A	214	23.193	11.814	66.177	1.00 30.44	AAAA
MOTA	1701	N	GLY	A	215	22.243	13.590		1.00 29.95	AAAA
ATOM	1702	CA	GLY	A	215	22.128	14.325	66.437	1.00 29.16	AAAA
ATOM	1703	С	GLY	A	215	23.222	15.379	66.582	1.00 28.54	AAAA
ATOM	1704	0	GLY	À	215	23.306	16.061	67.602	1.00 28.27	AAAA
MOTA	1705	Ŋ	TYR	A	216	24.063	15.521	65.561	1.00 27.35	AAAA
ATOM	1706	CA	TYR	A	216	25.150	16.497	65.616	1.00 27.08	AAAA
MOTA	1707	СВ	TYR			26.516	15.800	65.531	1.00 28.38	AAAA
ATOM	1708	CG			216	26.786	14.966	66.757	1.00 30.21	AAAA
ATOM	1709		TYR			26.138	13.735	66.955	1.00 29.81	AAAA
ATOM	1710		TYR			26.311	13.014	68.138	1.00 30.03	AAAA
ATOM	1711		TYR			27.619	15.450	67.774	1.00 29.71	AAAA
ATOM	1712		TYR			27.798	14.741	68.957	1.00 29.96	AAAA
	1713	CZ			216	27.143	13.528	69.138	1.00 30.84	AAAA
MCTA	1714	OH			216	27.143	12.859	70.332	1.00 31.14	AAAA
atom atom	1715	C			216	25.055	17.599	64.581	1.00 25.40	AAAA
	1715	o			216	26.046	18.240	64.243	1.00 26.38	AAAA
ATOM	7170	-		••		20.040	10.230			

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ATOM	1717	N	ASN A	21/	23.845	17.791	64.076	1.00 23.55	AAAA
MOTA	1718	CA	ASN A	217	23.549	18.830	63.119	1.00 21.52	AAAA
	1719	CB	ASN A		23.431	18.282	61.699	1.00 20.64	AAAA
MOTA									
ATOM	1720	CG	asn a		23.202	19.386	60.669	1.00 20.29	AAAA
MOTA	1721	OD1	ASN A	217	22.089	19.888	60.499	1.00 18.03	AAAA
ATOM	1722	MD2	ASN A	217	24.274	19.790	60.004	1.00 19.95	AAAA
							63.605		
ATOM	1723	С	ASN A		. 22.216	19.346		1.00 21.64	AAAA
MOTA	1724	.0	ASN A	217	21.263	18.576	63.757	1.00 20.34	AAAA
ATOM	1725	N	LEU A	218	22.165	20.647	63.873	1.00 22.22	AAAA
					20.960	21.282	64.388	1.00 22.03	AAAA
MOTA	1726	CA	LEU A						
MOTA	1727	CB	LEU A	218	21.195	21.711	65.840	1.00 20.97	AAAA
ATOM	1728	CG	LEU A	218	20.051	21.838	66.841	1.00 20.94	AAAA
			LEU A		20.513	22.744	67.936	1.00 20.31	AAAA
· ATOM	1729								
ATOM	1730	CD2	LEU A	218	18.818	22.412	66.227	1.00 21.27	AAAA
MOTA	1731	С	LEU A	218	20.669	22.513	63.547	1.00 22.70	AAAA
	1732	ō	LEU A		21.451	23.454	63.557	1.00 22.64	AAAA
MOTA									
MOTA	1733	N	asn a	219	19.564	22.491	62.808	1.00 24.00	AAAA
ATOM	1734	CA	ASN A	219	19.166	23.626	61.990	1.00 25.33	AAAA
	1735	CB	ASN A	219	18.656	23.190	60.614	1.00 26.94	AAAA
MOTA									
ATOM	1736	CG	ASN A	219	19.737	22.601	59.749	1.00 26.68	AAAA
MOTA	1737	OD1	. ASN A	219	20.812	23.169	59.626	1.00 28.06	AAAA
ATOM	1738	כחות	ASN A	219	19.446	21.471	59.117	1.00 26.26	AAAA
							62.710	1.00 25.69	AAAA
MOTA	1739	C	ASN A		18.046	24.345			
MOTA	1740	0	asn a	219	17.118	23.706	63.210	1.00 27.51	AAAA
ATOM	1741	N	ILE A	220	18.122	25.667	62.753	1.00 25.05	AAAA
		CA	ILE A		17.107	26.457	63.428	1.00 25.87	AAAA
MOTA	1742								
MOTA	1743	CB	ILE A	220	17.733	27.331	64.557	1.00 25.04	AAAA
MOTA	1744	CG2	ILE A	220	16.654	28.152	65.227	1.00 25.24	AAAA
ATOM	1745	CG1			18.460	26.447	65.584	1.00 24.07	AAAA
						25.502	66.378	1.00 22.28	AAAA
MOTA	1746		ILE A		17.557				
ATOM	1747	C	ILE A	220	16.430	27.370	62.414	1.00 26.20	AAAA
ATOM	1748	0	ILE A	220	16.801	28.534	62.265	1.00 25.35	AAAA
		N	PRO A		15,421	26.850	61.704	1.00 26.70	AAAA
MOTA	1749								
ATOM	1750	CD	PRO A			25.501	61.778	1.00 27.17	AAAA
ATOM	1751	CA	PRO A	221	14.706	27.640	60.703	1.00 27.67	AAAA
ATOM	1752	CB	PRO A	221	13.771	26.613	60.064	1.00 26.81	AAAA
		CG	PRO A		14.473	25.293	60.346	1.00 27.36	AAAA
MOTA	1753								
ATOM	1754	С	PRO A	221	13.944	28.763	61.390	1.00 28.61	AAAA
ATOM	1755	0	PRO A	221	13.218	28.515	62.363	1.00 29.91	AAAA
ATOM	1756	N	LEU A	222	14.100	29.990	60.900	1.00 28.15	AAAA
						31,117		1.00 28.48	AAAA
MOTA	1757	CA	LEU A		13.408				
ATOM	1758	CB	LEU A	222	14.431	32.041	62.191	1.00 28.69	AAAA
ATOM	1759	CG	LEU A	222	15.187	31.394	63.371	1.00 28.67	AAAA
	1760		LEU A		16.304	32.300	63.837	1.00 28.62	AAAA
MOTA							64.527	1.00 27.65	· AAAA
MOTA	1761	CDZ	LEU A		14.231	31.106			
MOTA	1762	С	LEU A	222	. 12 26	31.882	60.518	1.00 28.44	AAAA
MOTA	1763	0	LEU A	222	12.318	31.958	59.325	1.00 27.90	AAAA
		N	PRO A		. 1113	32.441	61.009	1.00 28.79	AAAA
ATOM	1764							1.00 29.20	AAAA
ATOM	1765	CD	PRO A		10.966	32.357	62.410		
MOTA	1766	CA	PRO A	223	10.437	33.202	60.227	1.00 29.36	AAAA
ATOM	1767	CB	PRO A	223	9.256	33.287	61.183	1.00 28.98	AAAA
						33.502	62.485	1.00 28.68	AAAA
ATOM	1768	CG	PRO A		9.965				
MOTA	1769	С	PRO A	223	10.890	34.585	59.753	1.00 30.15	AAAA
ATOM	1770	0	PRO A	223	11.864	35.152	60.253	1.00 30.18	AAAA
			LYS A	224	10.150	35.112	58.781	1.00 30.50	AAAA
MOTA	1771	N	nio W	224				1.00 29.92	
MOTA	1772	CA	LYS A	224	10.398	36.422	58.213		AAAA
MOTA	1773	CB	LYS A	224	9.491	36.661	57.008	1.00 30.57	AAAA
	1774	CG	LYS A	224	9.588	35.676	55.893	1.00 30.06	AAAA
MOTA			TWC *	224		36.087	54.798	1.00 30.91	AAAA
MOTA	1775	CD	LYS A	224	8.640				
ATOM	1776	CE	LYS A	224	8.575	35.051	53.705	1.00 32.15	AAAA
	1777	NZ	LYS A	224	7.628	35.476	52.648	1.00 32,75	AAAA
MOTA			LYS A	271	10.050	37.468	59.260	1.00 29.75	AAAA
ATOM	1778	Ç	DID A	224				1.00 29.84	
ATOM	1779	0	LYS A	224	9.308	37.193	60.196		AAAA
ATOM	1780	N	GLY A	225	10.555	38.678	59.079	1.00 29.39	AAAA
		CA	GLY A	225	10.261	39.730	60.031	1.00 29.87	AAAA
MOTA	1781		GDI A	225		39.447	61.415	1.00 29.85	AAAA
MOTA	1782	С	GLY A	443	10.809	23.44/	* .		-

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ATOM	1783	0	GLY A	225	10.371	40.051	62.392	1.00 29.85	2222
ATOM	1784	N	LEU A						AAAA
	-				11.775		61.499	1.00 29.50	AAAA
MOTA	1785	CA	LEU A		12.374	38.175	62.778	1.00 29.80	AAAA
MOTA	1786	CB	LEU A	226	13.513	37.170	62.570	1.00 28.81	AAAA
ATOM	1787	CG	LEU A	226	14.097		63.820	1.00 27,29	AAAA
ATOM	1788	CDI	LEU A		13.132				
		CDI	L DEU A	220			64.275	1.00 26.06	AAAA
ATOM	1789	CD2	LEU A		15.455	35.888	63.538	1.00 27.03	AAAA
ATOM	1790	С	LEU A	226	12.936	39,428	63.448	1.00 30.68	AAAA
ATOM	1791	0	LEU A		13.636		62.804	1.00 30.57	
ATOM	1792	N	ASN A						AAAA
					12.624		64.729	1.00 31.46	AAAA
ATOM	1793	CA	ASN A		. 13.139	40.769	65.469	1.00 32.06	AAAA
MOTA	1794	CB	asn a	227	12.012	41.507	66.217	1.00 31.74	AAAA
ATOM	1795	CG	ASN A	227	11.291		67.234	1.00 32.07	AÀAA
MOTA	. 1796		ASN A		11.914				
		NED 2	. ADI A	200			68.104	1.00 31.61	AAAA
MOTA	1797		ASN A		9.962		67.141	1.00 31.59	AAAA
MOTA	1798	С	ASN A	227	14.225	40.334	66.444-	1.00 32.45	AAAA
ATOM	1799	0	ASN A	227	14.413	39.140	66.688	1.00 32.78	AAAA
MOTA	1800	N	ASP A		14.943		67.002		
								1.00 33.32	AAAA
ATOM	1801	CA	ASP A		16.017		67.928	1.00 34.75	AAAA
MOTA	1802	CB	ASP A	228	16.508	42.233	68.654	1.00 36.77	AAAA
ATOM	1803	CG	ASP A	228	17.154	43.238	67.714	1.00 37.28	AAAA
ATOM	1804	OD1	ASP A		17.662		66.652	1.00 37.78	
			ASP A						AAAA
MOTA	1805				17.180		68.054	1.00 37.21	AAAA
ATOM	1806	С	ASP A		15.707	39.892	68.964	1.00 34.93	AAAA
MOTA	1807	Ο.	ASP A	228	16.448	38.919	69.056	1.00 36.92	AAAA
ATOM	1808	N	ASN A	229	14.635		69.741	1.00 33.90	AAAA
ATOM	1809	CA	ASN A		14.268		70.775		
			ASN A					1.00 33.01	AAAA
ATOM	1810	CB			12.965		71.455	1.00 33.79	AAAA
MOTA	1811	CG	ASN A		13.131	40.663	72.369	1.00 34.04	AAAA
ATOM -	1812	OD1	ASN A	229	13.783	40.564	73.405	1.00 34.25	AAAA
ATOM	1813	ND2	ASN A	229	12.550		71.988	1.00 34.05	AAAA
ATOM	1814	C	ASN A		14.114	37.656	70.276		
		ō						1.00 32.98	AAAA
ATOM	1815		ASN A		14.529	36.697		1.00 32.77	AAAA
ATOM	1816	N.	GLU A	230	13.496	37.523	69.108	1.00 32.02	AAAA
MOTA	1817	CA	GLU A	230	13.277	36.227	68.516	1.00 30.72	AAAA
ATOM	1818	CB	GLU A	230	12.399	36.375	67.272	1.00 31.38	AAAA
MOTA	1819	CG	GLU A		11.006				
						36.896	67.583	1.00 31.02	AAAA
ATOM	1820	CD	GLU A		10.175	37.187	66.350	1.00 31.52	AAAA
ATOM	1821		GLU A		10.644	37.970	65.497	1.00 31.89	AAAA
MOTA	1822	OE2	GLU A	230	9.047	36.655	66.241	1.00 31.04	AAAA
ATOM	1823	С	GLU A	230	14.628	35.622	68.180	1.00 30.79	AAAA
ATOM	1824	0	GLU A		14.905	34.465	68.512	1.00 31.05	
					•				AAAA
ATOM	1825	N	PHE · A		15.490	36.412	67.553	1.00 30.05	AAAA
MOTA	1826	CA	PHE A		16.811	35.920	67.191	1.00 28.94	AAAA
ATOM	1827	CB	PHE A	231	17.632	37.015	66.528	1.00 29.33	AAAA
ATOM	1828	CG	PHE A	231	18.949	36.537	65.972	1.00 28.79	AAAA
ATOM	1829	CD1	PHE A		18.982	35.585	64.957	1.00 28.93	
	1830								AAAA
ATOM			PHE A		20.152	37.067	66.436	1.00 28.55	AAAA
MOTA	1831		PHE A		20.195	35.160	64.397	1.00 28.32	AAAA
MOTA	1832		PHE A		21.376	36.657	65.888	1.00 28.97	AAAA
ATOM	1833	CZ	PHE A	231	21.397	35.695	64.860	1.00 28.81	AAAA
ATOM-	1834	č	PHE A		17.559	35.443	68.413		
								1.00 28.25	AAAA
ATOM	1835	0	PHE A		17.999	34.302	68.485	1.00 27.97	AAAA
MOTA	1836	N	LEU A	232	17.691	36.329	-69.384	1.00 27.93	AAAA
ATOM	1837	CA	LEU A	232	18.425	36.003	70.590	1.00 27.93	AAAA
MOTA	1838	CB	LEU A		18.521	37.234	71.484	1.00 28.16	
									AAAA
ATOM	1839	CG	LEU A		19.220	38.37 <i>9</i>	70.747	1.00 27.96	AAAA
ATOM	1840		LEU A		19.203	39.629	71.587	1.00 27.57	AAAA
ATOM	1841	CD2	LEU A	232	20.639	37.955	70.387	1.00 27.76	AAAA
ATOM	1842	C	LEU A		17.815	34.851	71.340	1.00 27.95	
		ō	LEU A						AAAA
ATOM	1843				18.526	34.061	71.941	1.00 27.92	AAAA
ATCM	1844	И	PHE A		16.495	34.758	71.298	1.00 28.81	AAAA
MOTE	1845	CA	PHE A	233	15.786	33.685	71.972	1.00 30.27	AAAA
ATOM	1846	CB	PHE A	233	14.278	33.837	71.745	1.00 31.51	AAAA
ATOM	1847	CG	PHE A		13.465	32.710	72.308	1.00 32.38	
									AAAA
MOTA	1848	CDT	PHE A	ددء	13.257	32.599	73.677	1.00 33.66	AAAA

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MOTA	1849	CD2	PHE	A	233		12.928	31.741	71.467	1.00	33.51	AAAA
ATOM	1850	CE1	PHE	A	233		12.518	31.537	74.201	1.00	35.10	AAAA
	1851		PHE									
MOTA							12.193	30.677	71.975	1.00	34.21	AAAA
MOTA	1852	CZ	PHE	A	233		11.986	30.572	73.344	1.00	35.23	AAAA
ATOM	1853	С	PHE	A	233		16.219	32.301	71.483		30.55	AAAA
	1854	ō			233							
- MOTA							16.438	31.391	72.280	1.00	30.65	AAAA
ATOM	1855	N	ALA	A	234		16.317	32.151	70.165	1.00	30.21	AAAA
MOTA	1856	CA	AT.A	Δ	234		16.698	30.892	69.549		28.97	
												AAAA
MOTA	1857	CB	ALA	A	234		16.398	30.942	68.065	1.00	30.40	AAAA
MOTA	1858	С	ALA	A	234		18.169	30.571	69.761	1.00	28.27	AAAA
ATOM	1859	0			234		18.564					
								29.401	69.830		26.6 <del>9</del>	AAAA
MOTA	1860	N	LEU	Α	.235		18.978	31.614	69.855	1.00	27.56	AAAA
MOTA	1861	CA	LEU	Α	235	•	20.402	31.427	70.055-	1.00	29.17	AAAA
MOTA	1862	CB	T ETT	7	235							
							21.126	32.767	69.989		29.04	AAAA
ATOM	1863	CG			235		22.527	32.757	69.378	1.00	28.54	AAAA
ATOM	1864	CD1	LEU	Α	235		23.350	33.837	70.058	1.00	2705	- AAAA
ATOM	1865		LEU				23.182					
								31.408	69.558		27.21	AAAA
ATOM	1866	С			235		20.637	30.799	71.429	1.00	30.99	AAAA
MOTA	1867	Ο.	LEU	Α	235		21.159	29.697	71.547	1.00	31.65	AAAA
ATOM	1868	N			236		20.242	31.514	72.471			
											31.88	AAAA
ATOM	1869	CA	GLU	A	236		20.409	31.042	73.838	1.00	32.99	AAAA
ATOM	1870	CB	GLU	A	236		19.689	31.990	74.790	1.00	34.63	AAAA
ATOM	1871		GLU				19.980	33.449	74.531			
											36.79	AAAA
MOTA	1872	CD			236		19.044	34.360	75.294	1.00	38.99	AAAA
ATOM	1873	OE1	GLU	Α	236		17.803	34.303	75.070	1.00	39.03	AAAA
ATOM	1874	OE2	GLU	Δ	236		19.559	35.132	76.126		41.56	
												AAAA
MOTA	1875	С			236		19.806	29.656	73.982	1.00	32.94	AAAA
MOTA	1876	0	GLU	Α	236		20.379	28.753	74.595	1.00	31.76	AAAA
ATOM	1877	N	LYS	Α	237		18.631	29.503	73.399		32.83	AAAA
	1878	CA										
ATOM					237		17.906	28.256	73.471		33.59	AAAA
MOTA	1879	CB	LYS	Α	237		16.504	28.506	72.942	1.00	35.00	AAAA
ATOM	1880	CG	LYS	Α	237		15.516	27.436	73.213	1.00	36.69	AAAA
ATOM	1881	CD			237		14.310	28.008	73.940		38.53	
												AAAA
MOTA	1882	CE	LYS	A	237		14.636	28.331	75.392	1.00	39.27	AAAA
ATOM	1883	NZ	LYS	Α	237		13.398	28.531	76.204	1.00	39.42	AAAA
ATOM	1884	С	LYS				18.619	27.129	72.707	1 00	33.14	
								•				AAAA
MOTA	1885	0	LYS				18.850	26.051	73.260	1.00	33.29	AAAA
ATOM	1886	N	SER	Α	238		18.985	27.374	71.452	1.00	32.13	AAAA
ATOM	1887	CA			238		19.671	26.345	70.685		31.25	AAAA
MOTA	1888	CB	SER				19.740	26.717	69.194	1.00	30.52	AAAA
MOTA	1889	OG	SER	Α	238		20.544	27.851	68.970	1.00	29.95	AAAA
MOTA	1890	С	SER	Α	238		21.075	26.064	71.236	1.00	31.21	AAAA
	1891	0										
MOTA			SER				21.556	24.929	71.169		30.06	AAAA
MOTA	1892	N	LEU	A	239	•	21.740	27.077	71.782	1.00	31.71	AAAA
ATOM .	1893	ÇA	LEU	Α	239		23.070	26.842	72.351	1.00	33.47	AAAA
MOTA	1894	CB	LEU				23.698	28.13C	72.900		31.25	
												AAAA
ATOM	1895	CG	LEU				23.988	29.300	71.97 <b>7</b>	1.00	29.80	AAAA
MOTA	1896		LEU				24.589	30.414	72.787	1.00	29.05	AAAA
ATOM	1897	CD2	LEU	Δ	239		24.919	28.903	70.872	1 00	29.36	AAAA
MOTA	1898	С	LEU				22.933	25.839	73.502		35.41	AAAA
ATOM	1899	0	LEU	Α	239		23.812	25.012	73.735	1.00	36.25	AAAA
MOTA	1900	N	GLU	A	240		21.816	25.906	74.213		37.34	AAAA
								_				
ATOM	1901	CA	GLU				21.594	25.005	75.331		39.39	AAAA
MOTA	1902	CB	GLU	A	240		20.281	25.361	76.017	1.00	41.90	AAAA
ATOM	1903	CG	GLU				20.040	24.610	77.308		45.52	AAAA
										_		
MOTA	1904	CD	GLU				19.665	25.552	78.432		47.80	AAAA
ATOM	1905	OEL	GLU	Α	240		18.670	26.295	78.274	1.00	49.70	AAAA
ATOM	1906	CE2	GLU				20.364	25.559	79.469		48.04	AAAA
MOTA	1907	C	GLD				21.583	23.555	74.875		38.80	AAAA
MOTA	1908	0	GLU	Α	240		22.224	22.700	75.478	1.00	37.85	AAAA
ATOM	1909	N	ILE	Α	241		20.847	23.293	73.804		39.66	AAAA
											_	
ATOM	1910	CA	ILE				20.751	21.955	73.223		40.81	AAAA
ATOM	1911	CB	ILE				19.912	21.994	71.917	1.00	41.10	AAAA
ATOM	1912	CG2	ILE	Α	241		19.850	20.621	71.287	1.00	40.88	AAAA
			ILE				18.502	22.514	72.220		41.45	AAAA
ATOM	1913											
ATOM	1914	CD1	ILE .	A	241		17.641	22.745	70.992	1.00	41.14	AAAA
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ATOM	1915	С	ILE A 241	22.159	21.424	72.893	1.00 41.66	AAAA
ATOM	1916	0	ILE A 241	22.445	20.229	73.045	1.00 42.10	AAAA
ATOM	1917	N	VAL A 242	23.026	22.324	72.432	1.00 41.42	AAAA
ATOM	1918	CA	VAL A 242	24.394	21.977	72.076	1.00 41.23	AAAA
ATOM	1919	CB	VAL A 242	25.089	23.146	71.351	1.00 40.40	AAAA
ATOM	1920	CG1	VAL A 242	26.556	22.850	71.171	1.00 39.25	AAAA
ATOM	1921	CG2	VAL A 242		23.384	70.004	1.00 39.79	AAAA
MOTA	1922	С	VAL A 242	25.228	21.604	73.298	1.00 42.67	AAAA
ATOM	1923	0	VAL A 242	25.882	20.562	73.316	1.00 41.83	AAAA
ATOM	1924	N	LYS A 243	25.198	22.456	74.318	1.00 44.41	AAAA
ATOM	1925	CA	LYS A 243	25.972	22.215	75.523	1.00 46.51	AAAA
MOTA	1926	CB	LYS A 243	25.797	23.363	76.522	1.00 47.29	AAAA
MOTA	1927	CG	LYS A 243	26.820	23.312	77.664	1.00 48.40	AAAA ·
ATOM	1928	CD	LYS A 243		24.248	78.823	1.00 48.88	AAAA
MOTA	1929	CE	LYS A 243	26.355	25.691	78.380	1.00 49.62	AAAA
ATOM	1930	NZ	LYS A 243	25.926	26.576	79.505	1.00 50.11	AAAA
ATOM	1931	C	LYS A 243		20.891	76.209	1.00 47.59	AAAA
MOTA	1932	0	LYS A 243		20.216	76.711	1.00 48.17	AAAA
MOTA	1933	N	GLU A 244		20.517	76.237	1.00 48.86	AAAA
MOTA	1934	CA	GLU A 244		19.262	76.877	1.00 50.82	AAAA
MOTA	1935	CB	GLU A 244		19.208	77.103	1.00 52.08	AAAA
ATOM	1936	CG	GLU A 244		20.405	77.829	1.00 53.82	AAAA
MOTA	1937	CD	GLU A 244		20.174	78.230	1.00 54.49	AAAA
ATOM	1938		GLU A 244		21.158	78.595	1.00 55.15	AAAA
MOTA	1939	OE2	GLU A 244		19.006	78.200	1.00 54.98	AAAA
MOTA	1940	С	GLU A 244		18.046	76.033	1.00 51.06	AAAA AAAA
ATOM	1941	0	GLU A 244		16.905	76.477	1.00 51.68 1.00 51.12	AAAA
MOTA	1942	N	VAL A 245		18.292	74.820 73.904	1.00 50.08	AAAA
MOTA	1943	CA	VAL A 245		17.212	72.677	1.00 50.22	AAAA
ATOM	1944	CB	VAL A 24		17.263	71.651	1.00 50.22	AAAA
ATOM	1945		VAL A 245		16.217 17.049	73.118	1.00 50.79	AAAA
ATOM	1946		VAL A 245		17.254	73.397	1.00 49.43	AAAA
MOTA	1947	C	VAL A 24!		16.250	72.917	1.00 48.65	AAAA
MOTA	1948	N O	PHE A 24		18.408	73.522	1.00 48.65	AAAA
ATOM	1949	CA	PHE A 240		18.552	72.982	1.00 47.97	AAAA
MOTA	1950 1951	CB	PHE A 24		19.212	71.607	1.00 46.45	AAAA
MOTA	1952	CG	PHE A 24		18.932	70.671	1.00 45.35	AAAA
ATOM ATOM	1953		PHE A 24		17.629	70.280	1.00 44.13	AAAA
MOTA	1954		PHE A 24		19.972	70.124	1.00 44.40	AAAA
MOTA	1955		PHE A 24		17.370	69.356	1.00 43.95	AAAA
MOTA	1956		PHE A 24		19.721	69.197	1.00 43.47	AAAA
ATOM	1957	CZ	PHE A 24		18.422	68.811	1.00 44.05	AAAA
ATOM	1958	C	PHE A 24		19.383	73.860	1.00 48.60	AAAA
MOTA	1959	o	PHE A 24	5 29.132	20.501	74.239	1.00 49.59	AAAA
ATOM	1960	N	GLU A 24	7 30.647	18.834	74.198	1.00 48.69	AAAA
ATOM .	1961	CA	GLU A 24	7 31.644	19.578	74.977	1.00 49.45	AAAA
ATOM	1962	CB	GLU A 24	7 32.174	18.768	76.178	1.00 51.91	AAAA
MOTA	1963	CG	GLU A 24	7 31.257	18.659	77.398	1.00 54.39	AAAA
MOTA	1964	CD	GLU A 24	7 29.986	17.845	77.146	1.00 57.34	AAAA
ATOM	1965	OE1	GLU A 24		18.315	76.393	1.00 58.48	AAAA
MOTA	1966	OE2	GLU A 24		16.725	77.702	1.00 57.95	AAAA
MOTA	1967	С	GLU A 24		19.903	74.024	1.00 47.39	AAAA
MOTA	1968	0	GLU A 24		19.119	73.872	1.00 46.65	AAAA
MOTA	1969	N	PRO A 24		21.070	73.371	1.00 46.25	AAAA
ATOM	1970	CD	PRO A 24		22.033	73.543	1.00 46.49	AAAA
MOTA	1971	CA	PRO A 24		21.614	72.411	1.00 45.44	AAAA
ATOM	1972	CB	PRO A 24		22.948	72.017	1.00 45.57	AAAA
ATOM	1973	CG	PRO A 24		22.661	72.178	1.00 46.28	AAAA
ATOM	1974	C	PRO A 24		21.814	72.880	1.00 44.29	AAAA
ATOM	1975	0	PRO A 24		22.370	73.947	1.00 44.57	AAAA
ATOM	1976	N	GLU A 24	36.100	21.364	72.059	1.00 42.21	AAAA
ATOM	1977	CA	GLU A 24		21.526	72.340	1.00 39.87	AAAA AAAA
MOTA	1978	CB	GLU A 24	9 38.344	20.460	71.625	1.00 39.58	AAAA
MOTA	1979	CG	GLU A 24		19.030	71.957	1.00 41.32	
ATOM	1980	CD	GLU A 24	9 38.825	18.007	71.241	1.00 40.96	AAAA •
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ATOM	1981	OE1	GLU	A 249	38.871	18.017	69.996	1.00 41.55	AAAA
MOTA	1982	OE2	GLU	A 249	39.462	17.187	71.926	1.00 42.49	AAAA
MOTA	1983	C	GLU	A 249	37.840	22:873	71.718	1.00 38.81	AAAA
MOTA	1984	0		A 249	38.715	23.617	72.181	1.00 38.27	AAAA
ATOM -	1985	N		A 250	37.109	23.160	70.644	1.00 36.60	AAAA
ATOM	1986	CA		A 250	37.242	24.402	69.890	1.00 34.20	AAAA
ATOM	1987	CB		A 250	38.379	24.321	68.862	1.00 33.73	AAAA
MOTA	1988			A 250	38.085	23.209	67.864	1.00 33.26	AAAA
MOTA	1989			A. 250	38.546	25.678	68.153	1.00 33.75	AAAA
MOTA	1990	C		A 250	35.945	24.617	69.130	1.00 31.98	AAAA
MOTA	1991	0		A 250	35.205	23.658	68.904	1.00 32.36	AAAA
MOTA	1992	N		A 251	35.657	25.863	68.760	1.00 28.65	AAAA
MOTA	1993	CA		A 251 A 251	34.449	26.150 26.442	67.991	1.00 26.49 1.00 24.32	AAAA
ATOM	1994 1995	CB		A 251	33.241		68.906	1.00 24.32	AAAA
ATOM		CG		A 251	33.193 32.771	27.853 28.931	69.465 68.668	1.00 22.36	AAAA
MOTA	1996 1997			A 251	32.791	30.234	69.151	1.00 21.29	AAAA AAAA
MOTA MOTA	1998			A 251	33.628	28.124	70.771	1.00 21.47	AAAA
MOTA	1999			A 251	33.651	29.425	71.265	1.00 20.80	AAAA
ATOM	2000	CZ		A 251	33.237	30.475	70.449	1.00 20.77	AAAA
ATOM	2001	OH		A 251	33.309	31.768	70.913	1.00 21.41	AAAA
ATOM	2002	C		A 251	34.691	27.345	67.092	1.00 24.59	AAAA
ATOM	2003	Ō		A 251	35.504	28.216	67.410	1.00 25.87	AAAA
ATOM	2004	N	LEU	A 252	33.984	27.374	65.970	1.00 22.49	AAAA
MOTA	2005	CA	LEU	A 252	34.082	28.482	65.045	1.00 20.96	AAAA
ATOM	2006	CB	LEU .	A 252	34.523	28.018	63.657	1.00 21.31	AAAA
MOTA	2007	CG		A 252	35.940	27.472	63.556	1.00 21.03	AAAA
MOTA	2008			A 252	35.947	26.028	63.977	1.00 22.16	AAAA
MOTA	2009			A 252	36.440	27.594	62.143	1.00 22.13	
MOTA	2010	C		A 252	32.731	29.159	64.959	1.00 19.60	AAAA
MOTA	2011	0		A 252	31.689	28.523		1.00 19.95 1.00 17.95	AAAA
MOTA	2012 2013	N CA		A 253 A 253	32.748 31.521	30.461 31.222	64.756 64.675	1.00 17.33	AAAA AAAA
MOTA	2013	CB		A 253	31.441	32.142	65.900	1.00 16.31	AAAA
ATOM ATOM	2015	CG		A 253	30.266	33.070	66.153	1.00 15.81	AAAA
ATOM	2016			A 253	28.990	32.267	66.377	1.00 14.74	AAAA
ATOM	2017			A 253	30.602		67.368	1.00 15.83	AAAA
ATOM	2018	С		A 253	31.564	32.035	63.386	1.00 16.60	AAAA
ATOM	2019	0	LEU .	A 253	32.548	32.722	63.132	1.00 16.40	AAAA
ATOM	2020	N	GLN .	A 254	30.526	31.936	62.557	1.00 15.88	AAAA
MOTA	2021	CA		A 254	30.507	32.716	61.328	1.00 16.27	AAAA
ATOM	2022	CB		A 254	30.045	31.881	60.121	1.00 15.88	AAAA
MOTA	2023	CG		A 254	28.587	32.048	59.734	1.00 18.52	AAAA
MOTA	2024	CD		A 254	28.380	32.935	58.519	1.00 17.54	AAAA
ATOM	2025	OE1		A 254 A 254	28.714 27.828	32.572	57.391 58.750	1.00 15.89	AAAA AAAA
MOTA	2026 2027	NE2 C		A 254	29.527	34.103 33.825	61.650	1 00 16.49	AAAA
ATOM ATOM	2028	Ö		A 254	28.450	33.571	62.198		AAAA
ATOM	2029	N	_	A 255	29.911	35.053	61.319	1.00 16.68	AAAA
ATOM	2030	CA		A 255	29.102	36.215	61.619	1.00 16.42	AAAA
ATOM	2031	CB		A 255	29.861	37.080	52.616	1.00 14.93	AAAA
ATOM		·CG		A 255	30.269	36.301		1.00 13.90	AAAA
MOTA	2033		LEU A		31.494	36.924	64.515	1.00 12.24	AAAA
ATOM	2034	CD2	LEU A	A 255	29.083	36.202	64.774	1.00 12.80	AAAA
ATOM	2035	С	LEU A	A 255	28.699	37.048	60.404	1.00 18.32	AAAA
MOTA	2036	0	LEU A	A 255	29.170	38.177	60.216	1.00 17.59	AAAA
MOTA	2037	N		A 256	27.813	36.482	59.588	1.00 19.75	AAAA
MOTA	2038	CA		A 256	27.322	37.188	58.422	1.00 20.77	AAAA
ATOM	2039	С	GLY A		26.422	38.302	58.927	1.00 21.73	AAAA
MOTA	2040	0	GLY 2		25.642	30.096	59.857	1.00 21.38	AAAA
MOTA	2041	N	THR A		26.528	39.485	58.325	1.00 22.82	AAAA
MOTA	2042	CA	THR A		25.721	40.622	58.746	1.00 23.85	AAAA
MOTA	2043	CB	THR A		26.460	41.968	58.549	1.00 23.99 1.00 25.54	AAAA AAAA
ATOM	2044	0G1	THR A	1.23/	26.729 27.780	42.169	57.153	1.00 23.34	AAAA
MOTA	2045		THR A		24.438	41.985	59.329 57.948	1.00 24.07	AAAA
ATOM	2046	C·	THR A	1 43/	44.438	40.691	21.348	1.00 44.31	Anna

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		_		_							•
ATOM	2047	0	THR	Α	257	23.692	41.672	58.048	1.00	25.84	AAAA
ATOM	2048	N	ACD	Δ	258	24.152	39.665	57.154	3 00	25.18	AAAA
ATOM	2049	CA	ASP	Α	258	22 <i>.</i> 935	39.753	56.379	1.00	26.18	AAAA
ATOM	2050	CB	ACD	Δ	258	22.950	38.830	55.149	1 00	25.52	AAAA
MOTA	2051	CG	ASP	A	258	23.211	37.392	55.494	1.00	26.33	AAAA
ATOM	2052	С	ASP	A	258	21.649	39:574	57.178	1 00	26.74	AAAA
ATOM	2053.	0	ĄSP	Α	258	20.571	39.823	56.643	1.00	26.57	AAAA
MOTA	2054	ODI	ASP	Δ	258	23.014	37.029	56.675	1 00	26.85	AAAA
					•						
ATOM	2055	OD2	ASP	Α	258	23.585	36:623	54.572	1.00	24.06	AAAA
ATOM	2056	N	PRO	Δ	259	21.727	39.114	58.449	1 00	26.95	AAAA
	_										
ATOM	2057	CD	PRO	Α	259 .	22.834	38.589	59.271	1.00	27.34	AAAA
ATOM	2058	CA	PRO	Α	259	20.467	38.976	59.190	1 00	27.05	AAAA
ATOM	2059	CB	PRO	A	259	20.886	38.186	60.425	1.00	26.38	AAAA
ATOM	2060	CĢ	PRO	Α	259	22.247	38.718	60.669	1.00	27.84	AAAA
MOTA	2061	С	PRO	A	259	19.914	40.365	59.543	1.00	27.32	AAAA
ATOM	2062	0	PRO	A	259	18.739	40.510	59.871-	1.00	27.29	AAAA
ATOM	2063	N			260	20.771	41.383	59.452	1.00	26.97	AAAA
ATOM	2064	CA	LEU	Α	260	20.389	42.763	59.752	1.00	26.74	AAAA
ATOM	2065	CB			260	21.621	43.680	59.680		27.21	AAAA
ATOM	2066	CG	LEU	Α	260	22.732	43.465	60.709	1.00	27.01	AAAA
ATOM	2067		LEU			23.889	44.380	60.408		25.51	AAAA
ATOM	2068	CD2	LEU	A	260 .	22.189	43.718	62.112	1.00	27.39	AAAA
	2069	c	TETT	70	260	19.295	43.351	58.865	3 00	26.47	
MOTA											AAAA
ATOM	2070	0	LEU	Α	260	19.278	43.137	57.649	1.00	26.72	AAAA
	2071	N	T.ETT	Z.	261	18.413	44.126	59.494	1 00	26.32	AAAA
MOTA	2072	CA	LEU	Α	261	17.283	44.808	58.846	1.00	27.20	AAAA
ATOM	2073	CB	UHU	A	261	16.732	45.885	59.780	1.00	28.71	AAAA
ATOM	2074	CG			261	15.644	46.789	59.19 <b>0</b>		29.24	AAAA
MOTA	2075	CD1	LEU	Α	261	14.433	45.954	58.883	1.00	29.44	AAAA
ATOM -	2076		LEU			15.284	47.906	60.162		29.72	AAAA
MOTA	2 <b>077</b>	С	LEU	A	261	17.506	45.454	57.473	1.00	27.90	AAAA
	2078	0	LEU	3	261	16.675	45.294	56.577	1 00	28.21	AAAA
ATOM											
ATOM	2079	И	GLU	Α	262	18.597	46.202	57.310	1.00	27.61	AAAA
ATOM	2080	CA	GI.II	A	262 .	18.887	46.877	56.043	1.00	26.92	AAAA
ATOM	2081	CB	GLU	A	262.	19.949	47.955	56.241		25.85	AAAA
MOTA	2082	CG	GLU	Α	262	19.549	49.119	57.117	1.00	25.36	AAAA
ATOM	2083	CD			262	19.552	48.787	58.580		25.78	AAAA
ATOM	2084	OE1	GLU	А	262	19.859	47.631	58.938	1.00	24.64	AAAA
	2085		GLU			19.255	49.694	59.381	1 00	25.82	AAAA
MOTA											
MOTA	2086	С	GLU	Α	262	19.346	45.995	54.882	1.00	28.79	AAAA
ATOM	2087	0	GLU	3	262	19.354	46.439	53.724	7 00	28.70	AAAA
ATOM	2088	N.	ASP	A.	263	19.743	44.758	55.179	T.00	29.57	AAAA
ATOM	2089	CA	ASP	A	263	20.230	43.853	54.145	1.00	28.99	AAAA
ATOM	2090	CB	ASP	A	263	21.160	42.802	54.760		27.89	AAAA
ATOM	2091	CG	ASP	Α	263	21.986	42.062	53.714	1.00	29.02	'AAAA
	2092		ASP			23.194	41.863	53.957		28.06	AAAA
ATOM											
ATOM	2093	ODZ	ASP	A	263	21.438	41.663	52.660	1.00	28.80	AAAA
ATOM	2094	С	ASP	Δ	263	19.066	43.197	53.431	1.00	29.73	. AAAA
MOTA	2095	0	ASP	Α	263	18.258	42.510	54.043	1.00	29.15	AAAA
MOTA	2096	N	TYR	Δ	264	19.002	43.416	52.122	1.00	31.25	AAAA
ATOM	2097	CA	TYR	A	264	17.925	42.88B ·	51.306		32.43	AAAA
MOTA	2098	CB	TYR	Α	264	17.913	43.558	49.938	1.00	34.53	AAAA
MOTA	2099	CG	TYR	A	264	17.627	45.038	49.997		38.21	AAAA
MOTA	2100	CD1	TYR	Α	264	18.664	45.968	· 49.983	1.00	39.87	AAAA
ATOM	2101	CEl	TYR			18.409	47.335	50.068		41.74	AAAA
ATOM	2102	CD2	TYR	A	264	16.316	45.511	50.103	1.00	40.10	AAAA
			TYR			16.044	46.877	50.191		41.50	AAAA
MOTA	2103										
ATOM	2104	cz	TYR	A	264	17.095	47.786	50.170	1.00	42.75°	AAAA
	2105	OH	TYR			16.838	49.147	50.231		44.65	AAAA
MOTA											
ATOM	2106	C	TYR	A	264	17.897	41.385	51.135	1.00	32.50	AAAA
ATOM	2107	0	TYR			16.819	40.816	50.968	1.00	3249	AAAA
ATOM	2108	И	LEU			19.064	40.740	51.171		32.64	AAAA
ATOM	2109	CA	LEU	Α	265	19.122	39.281	51.036	1.00	31.92	AAAA
MOTA	2110	CB	LEU			20.525	38.823	50.617		32.75	AAAA
ATOM	2111	CG	LEU	Α	265	20.808	39.010	49.127	1.00	32.95	AAAA
•											
MOTA	2112	CDI	LEU	M	203	22.213	38.588	48.771	1.00	31.59	AAAA

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MOTA	2113	CD2	LEU	Α	265	19.8	103 38.	166	48.361	1.00	34.62	AAAA
MOTA	2114	С	LEU	Δ	265	18.6	92 39	540	52.296	1 00	30.33	AAAA
ATOM	2115	0	LEU	А	265	19.0	24 37.	375	52.484	1.00	30.30	AAAA
ATOM	2116	N	SER	A	266	17.9	45 39	230	53.147	1 00	29.23	AAAA
MOTA	2117	CA	SER	А	266	17.4	34 38.	649	54.371	1.00	29.72	AAAA
ATOM -	2118	CB	SER	А	266	18.3	98 38.	894	55.519	1.00	32.09	AAAA
MOTA	2119	OG	SER	А	266	17.7	28 38.	810	56.771	1.00	33.43	AAAA
ATOM	2120	C	SER	Α	266	16.1	15 39.	290	54.698	1.00	28.71	ÀAAA
							-					
Mota	2121	0	SER	A	266	15.9			54.444	1.00	29.67	AAAA
ATOM	2122	N	LYS	Α	267	15.2	09 38.	517	55.276	1.00	27.82	AAAA
MOTA	2123	CA	LYS			13.9		045	55.654		27.56	AAAA
MOTA	2124	CB	LYS	Α	-267	12.8	21 38.	076	55.222	1.00	28.75	AAAA
	2125	CG	LYS	Δ	267	12.7	72 77	922	53.718-	1 00	29.67	AAAA
· ATOM												
MOTA	2126	CD	LYS	Α	267.	12.3	43 39.	223	53.053	1.00	30.13	AAAA
ATOM	2127	CE	LYS	Δ	267	12.3	03 39	036	51.546	1 00	31.86	AAAA
MOTA	2128	NZ	LYS	A	267	11.7	96 40.	252	50.843	T.00	33.92	-AAAA
ATOM	2129	С	LYS	Α	267	13.8	00 39.	327	57.152	1.00	27.18	· AAAA
MOTA	2130	0	LYS	A	267	12.7		591	57.665	1.00	27.18	AAAA
ATOM	2131	N	PHE	Α	268	14.9	44 39.	267	57.836	1.00	26.12	AAAA
		CA	PHE			15.0			59.271		25.72	
MOTA	2132											AAAA
MOTA	2133	CB	PHE	Α	268	16.2	72 38.	830	59.856	1.00	24.94	AAAA
ATOM.	2134	CG	PHE	Δ	268	16.1	67 37	334	59.896	1.00	25.07	AAAA
MOTA	2135	CDI	PHE	Α	268	17.2	71 36.	565	60.267	1.00	24.56	AAAA
ATOM	2136	CD2	PHE	A	268	14.9	55 36.	687	59.629	1.00	23.76	AAAA
MOTA	2137	CET	PHE	А	208	17.1	./4 35.	169	60.384		23.71	AAAA
MOTA	2138	CE2	PHE	Α	268	14.8	50 35.	303	59.739	1.00	23.86	AAAA
	2139	cz	PHE	Δ	268	15.9		542	60.121	1 00	23.68	AAAA
MOTA												
ATOM	2140	С	PHE	Ą	268	15.1	90 41.	030	59.513	1.00	25.77	AAAA
MOTA	2141	0	PHE	A	268	15.8	11 41.	734	58.726	1.00	25.81	AAAA
											26.02	
MOTA	2142	N	ASN	A	269	14.6		524	60.595			AAAA
MOTA	2143	CA	ASN	Α	269	14.7	18 42.	943	60.890	1.00	26.58	AAAA
		CB	ASN			13.3		584	61.058		25.47	AAAA
MOTA	2144											
MOTA	2145	CG	ASN	A	269	12.3	79 43.	252.	59.906	1.00	25.37	AAAA
MOTA	2146	ODI	ASN	Δ	269	12.7	61 43	260	58.734	1.00	23.82	AAAA
MOTA	2147	NUZ	ASN	A	269	11.1		985	60.245		24.03	AAAA
MOTA	2148	С	ASN	Α	269	15.5	40 43.	112	62.169	1.00	26.82	AAAA
	2149	Ō	ASN			15.0		715	63.150	1 00	27.98	AAAA
MOTA												
ATOM	2150	N	LEU	А	270	16.7	44 42.	559	62.149	1.00	26.07	AAAA
ATOM	2151	CA	LEU	Α	270	17.6	39 42	642	63.289	1.00	25.97	AAAA
											23.76	AAAA
ATOM	2152	CB	LEU			18.6		479	63.265			
MOTA	2153	CG	LEU	Α	270	18.0	48 40.	070	63.225	1.00	23.36	AAAA
	2154	CD1	LEU	Δ	270	19.1		090	63.710	1 00	21.90	AAAA
ATOM												
MOTA	2155	CD2	LEU	А	270	16.8	24 39.	971	64.122	1.00	22.05	AAAA
MOTA	2156	С	LEU	Α	270	18.4	20 43.	961	63.360	1.00	27.13	AAAA
			LEU					750	62.399		25.99	AAAA
MOTA	2157	0				18.4						
MOTA	2158	N	SER	A	271	19.0	38 44.	176	64.517	1.00	27.97	: AAAA
ATOM	2159	CA	SER	Δ	271	19.8	32 45	370	64.767	1.00	27.95	AAAA
ATOM	2160	CB	SER	A	271	19.2	35 46.	137	65.943	1.00	27.32	AAAA
ATCM	2161	OG	SER	Α	271	19.1	84 45.	297	67.089	1.00	27.90	AAAA
											28.15	AAAA
MOTA	2162	С	SER			21.2		987	65.084			
MOTA	2163	0	SER	Α	271	21.5	74 43.	832	65.401	1.00	26.99	AAAA
	2164	N	ASN			22.1			64.979	. 1.00	28 96	, AAAA
ATOM												
MOTA	2165	CA	ASN	А	272	23.5	90 45.	861	65.266	1.00	29.54	AAAA
	2166	CB	ASN	A	272	24.2	47 47.	243	65.223	1.00	30.96	AAAA
ATOM												
ATOM	2167	CG	ASN	A	212	24.6	47 47.	640	63.850		31.20	AAAA
ATOM	2168	OD1	ASN	Α	272	24.9	60 48.	794	63.594	1.00	31.73	AAAA
											31.93	AAAA
ATOM	2169		ASN			24.6			62.948			
ATOM	2170	С	ASN	A	272	23.8	21 45.	309	66.645	1.00、	29.63	AAAA
		ō			_	24.5			66.843		29.85	AAAA
MOTA	2171		ASN									
MOTA	2172	N	VAL	A	273	23.1	80 45.	<b>959</b>	67.600	1.00	29.77	AAAA
	2173	CA	VAL			23.2	90 45.	602	68.994	1.00	30.89	AAAA
MOTA												
MOTA	2174	CB	VAL	A	213	22.4			69.816		31.61	AAAA
MOTA	2175	CG1	VAL	Α	273	22.7	16 46.	403	71.293	1.00	33.17	AAAA
						22.7		-			31.82	AAAA
MOTA	2176		VAL					998	69.372			
MOTA	2177	С	VAL	A	273	22.8	83 44.	144	69.266		30.74	AAAA
	2178	Ō	VAL			23.5			70.022		31.23	AAAA .
MOTA	٠,٠٠	9	A LTE			45.5						

SUBSTITUTE SHEET (RULE 26)



MOTA	2179	N	ALA A	274	21.785	43.706	68.659	1.00 30.25	AAAA
ATOM	2180	CA	ALA A		21.327	42.333	68.840	1.00 29.87	AAAA
		CB	ALA A		20.005	42.119	68.112	1.00 29.64	AAAA
MOTA	2181							•	
ATOM	2182	С	ALA A		22.395	41.438	68.247	1.00 29.35	AAAA
MOTA	2183	0	ALA A		22.707	40.373	68.778	1.00 29.18	AAAA
MOTA	2184	N	PHE A	275	22.946	41.893	67.127	1.00 29.30	AAAA
ATOM	2185	CA	PHE A	275	23.991	41.170	66.428	1.00 28.91	AAAA
	2186	CB	PHE A		24.375	41.909	65.150	1.00 28.77	AAAA
ATOM									
MOTA	2187	CG	PHE A		25.354	41.170	64.308	1.00 28.08	AAAA
MOTA	2188		PHE A		25.015	39.954	63.740	1.00 28.92	AAAA
MOTA	2189	CD2	PHE A	275	26.621	41.684	64.077	1.00 29.48	AAAA
MOTA	2190	CE1	PHE A	275	25.928	39.259	62.945	1.00 29.20	AAAA
ATOM	2191	CE2	PHE A	275	27.546	40.988	63.279	1.00 29.24	AAAA
	2192	CZ	PHE A		27.193	39.779	62.716	1.00 28.30	AAAA
MOTA					25.196			1.00 27.64	
MOTA	2193	C	PHE A			41.058	67.351		AAAA
ATOM	2194	0	PHE A		25.728	39.975	67.558	1.00 28.65	AAAA
MOTA	2195	N	LEU A	276	25.606	42.189	67.902	1.00 26.81	AAAA
ATOM	2196	CA	LEU A	276	26.732	42.260	68.831	1.00 27.38	AAAA
ATOM	2197	CB	LEU A	276	26.878	43.700	69.353	1.00 27.53	AAAA
ATOM	2198	CG	LEU A	276	28.202	44.213	69.928	1.00 26.37	AAAA
	2199		LEU A		27.923	45.488	70.721	1.00 25.71	AAAA
MOTA							70.827	1.00 27.06	
MOTA	2200		LEU A		28.842	43.189			AAAA
MOTA	2201	C	LEU A		26.486	41.317	70.021	1.00 26.49	AAAA
MOTA	2202	0	LEU A	276	27.387	40.603	70.471	1.00 25.26	AAAA
ATOM .	2203	N	LYS A	277	25.257	41.322	70.524	1.00 27.46	AAAA
MOTA	2204	CA	LYS A	2 <b>7</b> 7	24.894	40.468	71.642	1.00 28.63	AAAA
ATOM	2205	CB	LYS A		23.542	40.862	72.223	1.00 30.63	AAAA
ATOM	2206	CG	LYS A		23.590	42.029	73.153	1.00 33.14	AAAA
		CD	LYS A		22.599	41.791	74.268	1.00 34.94	AAAA
ATOM	2207							1.00 36.17	AAAA
ATOM	2208	CE	LYS A		22.964	40.519	75.029		
ATOM	2209	NZ	LYS A		21.979	40.194	76.104	1.00 38.64	AAAA
ATOM	2210	С	LYS A	277	24.846	38.997	71.297	1.00 28.53	AAAA
ATOM	2211	0	LYS A	27 <b>7</b>	25.118	38.152	72.146	1.00 28.45	AAAA
ATOM	2212	N	ALA A	278	24.466	38.681	70.064	1.00 28.47	AAAA
MOTA	2213	CA	ALA A		24.404	37.280	69.656	1.00 27.66	AAAA
	2214	CB	ALA A		23.941	37.181	68.201	1.00 26.40	AAAA
ATOM			ALA A		25.833	36.754	69.820	1.00 26.63	AAAA
ATOM	2215	C							
MOTA	2216	0	ALA A		26.081	35.644	70.317	1.00 25.19	AAAA
MOTA	2217	N	PHE A		26.764	37.616	69.427	1.00 26.50	AAAA
MOTA	2218	CA	PHE A		28.181	37.345	69.481	1.00 25.83	AAAA
ATOM	2219	CB	PHE A	279	28.934	38.521	68.869.	1.00 26.35	AAAA
ATOM	2220	CG	PHE A	279	30.413	38.319	68.796	1.00 27.92	AAAA
ATOM	2221	CD1	PHE A	279	30.949	37.256	68.072	1.00 28.58	AAAA
ATOM	2222		PHE A		31.280	39.201	69.434	1.00 28.33	AAAA
	2223		PHE A		32.330	37.078	67.983	1.00 28.22	AAAA
MOTA						39.030	69.349	1.00 28.11	AAAA
MOTA	2224		PHE A		32.666			1.00 28.21	
MOTA	2225	cz	PHE A		33.185	37.968	68.622		AAAA
ATOM	2226	C	PHE A	279	28.665	37.118	70.901	1.00 25.47	AAAA
MOTA	2227	0	PHE A	279	29.284	36.091	71.202	1.00 24.32	AAAA
ATOM	2228	N	ASN A	280	28.382	38.075	71.778	1.00 25.12	AAAA
	2229		ASN A	280		37.944	73.147	1.00 25.05	AAAA
	2230	CB	ASN A		28.708	39.269	73.887	1.00 24.42	AAAA
ATOM						40.300	73.364	1.00 24.56	AAAA
ATOM	2231	CG	ASN A		29.683				
MOTA	2232		ASN A		30.841	39.980	73.080	1.00 23.24	AAAA
ATOM	2233	ND2	ASN A	280	29.233	41.543	73.249	1.00 24.59	AAAA
ATOM	2234	С	ASN A	280	28.213	36.814	73.925	1.00 24.79	AAAA
ATOM	2235	0	ASN A		28.828	36.272	74.825	1.00 24.96	AAAA
	2236	N	ILE A		26.998	36.444	73.565	1.00 24.87	AAAA
ATOM		CA	ILE A		26.332	35.337	74.220	1.00 24.80	AAAA
ATOM	2237					35.252	73.780	1.00 24.40	AAAA
MOTA	2238	CB	ILE A		24.866				
MOTA	2239		ILE A		24.297	33.907	74.124	1.00 25.03	AAAA
ATOM	2240		ILE A		24.076	36.386	74.424	1.00 24.70	AAAA
ATOM	2241	CD1	ILE A		22.613	36.379	74.069	1.00 26.49	AAAA
ATOM	2242	C	ILE A	281	27.044	34.027	73.884	1.00 25.21	AAAA
ATOM	2243	0	ILE A		27.220	33.170	74.750	1.00 24.97	AAAA
ATCM	2244	N	VAL A		27.440	33.866	72.620	1.00 25.98	AAAA
-A 11.TPI	4477								

MOTA	2245	CA	VAL	Α	282	28.150	32.656	72.193	1.00 2	25.15	AAAA
ATOM	2246	CB	VAL	Α	282	28.451	32.666	70.677	1.00 2	23.83	AAAA
MOTA	2247		VAL			29.315	31.470	70.311	1.00	23.58	AAAA
	2248		VAL			27.173	32.633	69.899	1.00 2		AAAA
MOTA	2249	C	VAL			29.478	32.553	72.936	1.00		AAAA
MOTA	2250	ō	VAL			29.928	31.457	73.275	1.00		AAAA
MOTA			ARG			30.100	33.702	73.176	1.00		AAAA
MOTA	2251	N				31.372	33.760	73.885	1.00		AAAA
MOTA	2252	CA	ARG				35.131	73.684	1.00		
ATOM	2253	CB	ARG			32.027					AAAA
MOTA	2254	CG	ARG			32.364	35.440	72.240	1.00		AAAA
ATOM	2255	CD.	ARG			32.821	36.862	72.098	1.00		AAAA
ATOM	2256	NE	ARG			34.035	37.116	72.854	1.00		AAAA
ATOM.	2257	CZ	ARG			34.514	38.327	73.091	1.00		AAAA
MOTA	2258	NH1	ARG	A	283	33.873	39.384	72.626	1.00		AAAA
MOTA	2259	NH2	ARG	Α	283	35.622	38.484	73.798	1.00		AAAA
ATOM	2260	С	ARG	Α	283	31.183	33.494	75.376	1.00		AAAA
ATOM	2261	0	ARG	A	283	32.086	32.981	76.027	1.00	30.68	AAAA
ATOM	2262	N	GLU	Α	284	30.014	33.842	75.911	1.00	32.71	AAAA
ATOM	2263	CA	GLU	Α	284	29.735	33.623	77.323	1.00	35.53	AAAA
ATOM	2264	CB	GLU	Α	284	28.482	34.391	77.751	1.00	37.39	AAAA
ATOM	2265	CG	GLU			28.538	35.854	77.392	1.00	41.73	AAAA
ATOM	2266	CD	GLU			27.272	36.631	77.754	1.00	45.27	AAAA
ATOM	2267		GLU			26.151	36.078	77.610	1.00	46.66	AAAA
ATOM	2268	OE2	GLU			27.405	37.817	78.148	1.00	46.94	AAAA
	2269	C			284	29.524	32.133	77.564	1.00	36.25	AAAA
MOTA	2270	ō.	GLU			29.920	31.593	78.601	1.00		AAAA
MOTA	2271	N	VAL			28.916	31.464	76.591	1.00		AAAA
MOTA		CA			285	28.637	30.041	76.708	1.00		AAAA
ATOM	2272	CB	VAL			27.505	29.619		1.00		AAAA
MOTA	2273		VAL			27.201	28.137	75.888	1.00		AAAA
ATOM	2274					26.254	30.457	76.001	1.00		AAAA
ATOM	2275				285	29.847	29.149	76.456	1.00		AAAA
ATOM	2276	C			285	30.140	28.262	77.257	1.00		AAAA
MOTA	2277	0	VAL				29.389	75.364		32.34	AAAA
ATOM	2278	N			286	30.568 31.706	28.535	75.036		29.92	AAAA
ATOM	2279	CA			286	31.533	27.960	73.635		29.77	AAAA
MOTA	2280	CB			286 286	30.267	27.179	73.444		28.64	AAAA
ATOM	2281	CG	PHE			29.152	27.772	72.863		28.75	AAAA
ATOM	2282		PHE			30.197	25.837	73.827		28.55	AAAA
ATOM	2283		PHE			27.983	27.039	72.660		29.04	AAAA
	. 2284 2285		PHE			29.037	25.095	73.629		28.19	AAAA
ATOM		CZ			286	27.929	25.694	73.045		28.73	AAAA
MOTA	2286	C			286	33.106	29.113	75.132		29.13	AAAA
ATOM	2287	ō	PHE			34.073	28.436	74.760		28.54	AAAA
MOTA	2288		GLY			33.224	30.341	75.637		28.42	AAAA
MOTA	2289	N	GLY			34.525	30.987	75.744		27.07	AAAA
ATOM	2290	CA				34.932	31.611	74.419		26.64	AAAA .
ATOM	2291	C	GLY				32.042	73.649		27.13	AAAA
MOTA	2292	0	GLY			34.088		74.146		27.20	AAAA
Mota	2293	N	GLU			36.227	31.665	72.900		27.52	AAAA
MOTA	2294	CA	GLU			36.719	32.238	73.108		28.18	AAAA
Mota	2295	CB			288	38.073	32.923	73.108		28.88	AAAA
MOTA	2296	CG	GLU			38.036	34.177	73.279		29.58	
MOTA	2297	CD	GLU			37.329	35.330	72.243		29.94	AAAA AAAA
MOTA	2298		GLU			37.807	35.813				
ATOM	2299	OE2	GLU			36.281	35,761	73.782		31.89	AAAA
ATOM	2300	С	GLU			36.877	31.158	71.843		27.44	AAAA
ATOM	2301	0	GLU			37.169	30.007	72.162		27.87	AAAA
MOTA	2302	N	GLY			36.663	31.547	70.589		26.55	AAAA
MOTA	2303	CA	GLY			36.795	30.638	69.466		25.25	AAAA
ATOM	2304	C	GLY			37.285	31.414	68.254		24.55	AAAA
ATOM	2305	0	GLY	А	289	37.635	32.586	68.369		24.40	AAAA
MOTA	2306	N	VAL	A	290	37.320	30.765	67.095		24.04	AAAA
MOTA	2307	CA	VAL	Α	290	37.756	31.407	65.863		23.76	AAAA
ATOM	2308	CB	VAL			38.288	30.346	64.867		24.94	AAAA
ATOM	2309	CG1	VAL	A	290	38.835	31.012	63.596		22.73	AAAA
MOTA	2310	CG2	VAL	A	290	39.375	29.506	65.555	1.00	24.74	AAAA
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ATOM	2311	С	VAL	Α	290	36.536	32.122	65.277	1.00	23.90	<b>AAAA</b>
ATOM	2312	0	VAL	А	290	35.497	31.502	65.100	1.00	25.15	AAAA
								64.976	1.00		AAAA
ATOM	2313	N	TYR			36.662	33.415				
ATOM	2314	CA	TYR	Α	291	35.544	34.211	64.446	1.00	21.41	AAAA
	2315	CB	TYR	Δ	291	35.472	35.540	65.193	1.00	20.57	AAAA
MOTA											
MOTA	2316	CG	TYR			35.511	35:346	66.677	1.00		AAAA
ATOM	2317	CD1	TYR	Α	291	36.596	35.782	67.432	1.00	20.86	AAAA
	•		TYR			36.677	35.513	68.793	1.00	21 47	AAAA
MOTA	2318										
ATOM	2319	CD2	TYR	Α	.291	34.509	34:647	67.318	1.00	20.90	AAAA
ATOM	2320	CE2	TYR	Α	291	34.579	34.372	68.675	1.00	21.90	AAAA
								69.403	1.00		AAAA
ATOM	2321	CZ	TYR			35.661	34.800				
MOTA	2322	OH	TYR	Α	291	35.737	34.469	70.730	1.00	23.75	AAAA
ATOM	2323	С	ጥV₽	Δ	291	35,607	34.483	62.946	1.00	21.25	AAAA
								62.451	1.00		AAAA
MOTA	2324	0			291	36.573	35.077				
ATOM	2325	N	LEU	А	292	34.557	34.084	62.231	1.00	20.92	AAAA
	2326	CA	LEU	Δ	292	34.518	34.260	60.779~	1.00	20.92	AAAA
ATOM											
ATOM	2327	CB	LEU	A	292 .	34.235	32.916	60.080	1.00		AAAA
ATOM	2328	CG	LEU	A	292	35.104	31.688	60.399	1.00	17.31	AAAA
	2329	CD1	LEU	Δ.	292	34.685	30.515	59.528	1.00	16.05	AAAA
MOTA											
ATOM	2330	CD2	LEU			36.552	32.000	60.163	1.00		AAAA
MOTA	2331	С	LEU	Α	292	33.515	35.288	60.283	1.00	21.12	AAAA
		0			292	32.652	35.741	61.020	1.00	20 70	AAAA
MOTA	2332										
ATOM	2333	N	GLY	А	293	33.660	35.660	59.017		21.74	AAAA
ATOM	2334	CA	GLY	Α	293	32.752	36.612	58.410	1.00	21.48	AAAA
		C			293	31.612	35.856	57.770	1 00	21.65	AAAA
ATOM	2335										
ATOM	2336	0	GLY	Α	293	31.237	34.790	58.235		22.25	AAAA
ATOM	2337	N	GLY	Α	294	31.060	36.392	56.691	1.00	22.66	AAAA
						29.957	35.714	56.034	3 00	23.61	AAAA
MOTA	2338	CA			294						
ATOM	2339	C	GĻY	Α	294	29.180	36.653	55.146		24.56	AAAA
ATOM -	2340	0	GLY	Α	294	29.679	37.727	54.790	1.00	25.54	AAAA
					295	27.956	36.265	54.794		24.06	AAAA
MOTA	2341	N									
ATOM	2342	CA	GLY	Α	295	27.139	37.093	53.927	1.00	22.78	AAAA
ATOM	2343	С	GI.Y	Α	295	26.902	38.479	54.483	1.00	23.11	AAAA
						26.870	38.676	55.696		22.87	AAAA
ATOM	2344	O _.			295						
ATOM	2345	N	GLY	Α	296	26.733	39.442	53.584		22.78	AAAA
ATOM	2346	CA	GLY	Α	296	26.497	40.813	53.993	1.00	23.44	AAAA
					296	26.471	41.618	52.718	1.00	23.72	AAAA
MOTA	2347	C									
ATOM	2348	0	GLY	A	296	27.474	41.661	52.004		23.73	AAAA
ATOM	2349	N	TYR	А	297	25,356	42.280	52.425	1.00	23.41	AAAA
					297	25.282	42.991	51.163	1.00	22.71	AAAA
MOTA	2350	CA									
ATOM	2351	CB	TYR	Α	297	24.252	42.294	50.296		21.55	AAAA
MOTA	2352	CG	TYR	Α	297	24.496	40.809	50.317	1.00	21.93	AAAA
			TYR			24.036	40.016	51.375	1.00	20.95	AAAA
MOTA	2353										
MOTA	2354		TYR			24.400	38.678	51.481		21.59	AAAA
ATOM	2355	CD2	TYR	Α	297	25.320	40.217	49.358	1.00	21.71	AAAA
			TYR			25.688	38.900	49.451	1 00	21.99	AAAA
MOTA	2356										
MOTA	2357	CZ				25.242	38.127	50.511		22.18	AAAA
ATOM	2358	OH	TYR	A	297	25.721	36.841	50.615	1.00	21.35	AAAA
		c	מעיווי	Δ	297	25.042	44.485	51.225	1.00	22.90	AAAA
ATOM	2359							50.203		23.17	AAAA
ATOM	2360	0			297	25.106	45.172				
ATOM	2361	N	HIS	A	298	24.772	44.989	52.417		22.47	AAAA
		CA	HIS			24.572	46.415	52.566	1.00	24.27	AAAA
MOTA	2362		1113		200						AAAA
ATOM	2363	CB	HIS			23.468	46.726	53.556		23.17	
ATOM	2364	CG	HIS	Α	298	23.097	48.166	. 53.572	1.00	23.20	AAAA
	2365		HIS			23.588	49.201	54.287	1.00	24.25	AAAA
ATOM		- LUC	****		200			52.680		23.14	AAAA
ATOM	2366	NDI	HIS	A	230	22.199	48.708				
ATOM	2367	CEl	HIS	A	298	22.151	50.017	52.848		23.31	AAAA
		MEG	HIS	A	298	22.986	50.342	53.814	1.00	23.62	AAAA
MOTA	2368									25.17	AAAA
MOTA	2369	С			298	25.886	46.976	53.106			
MOTA	2370	0	HIS	A	<b>298</b>	26.282	46.687	54.239	1.00	24.47	AAAA
					299	26.563	47.818	52.316	1.00	26.37	AAAA
MOTA	2371	N								27.01	AAAA
ATOM	2372	CD			299	26.178	48.372	51.006			
ATOM	2373	CA	PRO	A	299	27.840	48.394	52.752		27.31	AAAA
		CB			299	28.156	49.383	51.630	1.00	27.04	AAAA
MOTA	2374							51.120		27.57	AAAA
MOTA	2375	CG			299	26.743	49.764				
ATOM	2376	С	PRO	Α	299	27.824	49.037	54.149	1.00	27.77	AAAA
	23,0							•			•
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ATOM	2377	0	PRO	Α	299		28.755	48.826	54.939	1.00.	28.04	AAAA
ATOM	2378	N			300		26.769	49.794	54.452		27.04	AAAA
							26.629	50.477	55.740		27.59	AAAA
ATOM	2379	CA			300						-	
MOTA	2380	CB			300		25.425	51.437	55.700		30.57	AAAA
ATOM	2381	CG			300		25.516	52.599	54.718		32.91	AAAA
ATOM -	2382	CD1	TYR	Α	300		26.181	52.464	53.491	1.00	33.45	AAAA
MOTA	2383	CE1	TYR	Α	300		26.160	53.487	52.538	1.00	33.91	AAAA
ATOM	2384				300		24.837	53.801	54.969		34.19	AAAA
											34.64	
MOŢA	2385				300		24.809	54.830				AAAA
ATOM	2386	CZ	TYR	A	300		25.468	54.657	52.807		34.56	AAAA
ATOM	2387	OH	TYR	A	300		25.389	55.630	51.844	1.00	36.05	AAAA
MOTA	2388	С	TYR	Α	300		26.454	49.538	56.936	1.00	26.48	AAAA
ATOM	2389	ō			300		27.073	49.726	57.979 -	1.00	25.81	AAAA
		N			301		25.581	48.547	56.791		25.41	AAAA
MOTA	2390			-								
MOTA	2391	CA			301		25.328	47.606	57.865		24.64	AAAA
ATOM	2392	CB	ALA	A	301		24.164	46.731	57.511		25.32	-AAAA
ATOM	2393	C	ALA	A	301		26.568	46.775	58.067	1.00	25.53	AAAA
ATOM	2394	0	ALA	Α	301		27.030	46.567	59.194	1.00	26.39	AAAA
ATOM ·		И			302		27.108	46.304	56.950	1.00	25.83	AAAA
	2396	CÁ			302		28.323	45.500	56.926		26.32	AAAA
MOTA											27.38	AAAA
ATOM	2397	CB			302		28.782	45.378	55.479			
MOTA	2398	CG			302		30.081	44.723	55.024		28.18	AAAA
ATOM	2399	CD1	LEU	Α	302		30.119	44.840	53.502	1.00	29.32	AAAA
ATOM	2400	CD2	LEU	Α	302		31.296	45.389	55.613	1.00	27.38	AAAA
ATOM	2401	С	LEU	Α	302		29.398	46.187	57.764	1.00	26.41	AAAA
ATOM	2402	ō			302		29.874	45.648	58.755		26.62	AAAA
		N			303		29.756	47.397	57.353		26.50	AAAA
MOTA	2403										25.92	
ATOM	2404	CA			303			48.176	58.022			AAAA
MOTA	2405	CB			303		31.001	49.475	57.277		25.24	AAAA
MOTA	2406	С	ALA	Α	303		30.490	48.464	59.487		26.03	AAAA
ATOM	2407	0	ALA	. A	303	•	31.325	48.175	60.340	1.00	26.95	AAAA
MOTA	2408	N	ARG	Α	304		29.322	49.028	59.792	1.00	25.29	AAAA
MOTA	2409	CA			304		28.999	49.353	61.179	1.00	23.46	AAAA
	2410	CB			304		27.641	50.059	61.291		23.78	AAAA
MOTA			-			•			60.629		24.59	AAAA
MOTA	2411	-CG			304		27.553	51.451				
ATOM	2412	CD			304		26.302	52.223	61.091		25.85	AAAA
MOTA	2413	NE	ARG	A	304		25.067	51.465	60.869		27.54	AAAA
ATOM	2414	CZ	ARG	Α	304		23.978	51.547	61.637	1.00	28.36	AAAA
ATOM	2415	NH1	ARG	A	304		23.957	52.362	62.695	1.00	26.48	AAAA
ATOM	2416				304		22.910	50.794	61.358	1.00	28.45	AAAA
	2417	C			304		28.991	48.118	62.053		23.18	AAAA
ATOM					304		29.591	48.099	63.135		22.26	AAAA
MOTA	2418	0							61.560		23.20	AAAA
MOTA	2419	N			305		28.330	47.075				
ATOM	2420	CA			305		28.200	45.817	62.292		22.33	AAAA
ATOM	2421	CB	ALA	A.	305		27.319	44.866	61.516		22.17	AAAA.
MOTA	2422	С	ALA	Α	305		29.516	45.137	62.621	1.00	22.27	AAAA
ATOM	2423	0			305		29.763	44.757	63.760	1.00	22.48	AAAA
	2424	N			306		30.366	44.969	61.620		22.57	AAAA
ATOM					306			44.307	61.861		21.28	AAAA
ATOM	2425	CA					31.634					
ATQM	2426	CB			306		32.279	43.885	60.553		21.07	AAAA
ATOM	2427	ĊG			306		31.703	42.618	60.004		20.75	AAAA
ATOM	2428	CD2	TRP	Α	306		31.886	42.103	58.683	1.00	19.54	AAAA
ATOM	2429	CE2	TRP	A	306		31.352	40.795	58.668	1.00	19.18	AAAA
	2430		TRP				32.456	42.616	57.510		19.59	AAAA
ATOM											20.51	AAAA
ATOM	2431		TRP				31.071	41.632	60.713.			
ATOM	2432		TRP				30.864	40.537	59.922		19.74	
MOTA	2433	CZ2	TRP	Α	306		31.368	39.990	57.524		19.18	AAAA
ATOM	2434	CZ3.	TRP	A	306		32.474	41.810	56.367	1.00	18.98	AAAA
ATOM	2435	CH2	TRP	A	306		31:933	40.513	56.388	1.00	19.21	AAAA
		C	פקיד	Δ	306		32.571	45.159	62.674		20.80	AAAA
ATOM	2436								63.341		20.55	AAAA.
MOTA	2437	0			306		33.459	44.630				
ATOM	2438	N	THR	A	307		32.373	46.475	62.614		20.17	AAAA
MOTA	2439	CA	THR	Α	307		33.175	47.399	63.407		20.54	AAAA
ATOM	2440	CB	THR	A	307		32.861	48.881	63.045		21.09	AAAA
ATOM	2441	OG1	THR	Α	307		33.329	49.159	61.718	1.00	21.25	AAAA
	2442	CCS	THR	Δ	307		33.523	49.839	64.030		20.09	AAAA
MOTA	6334	C02	1111/	••			JJ.J-J	20.000				•



ATOM	2443	С	THR A 307	32.853	47.135	64.893	1.00 20.88	AAAA
ATOM	2444	ō	THR A 307	33.738	47.175	65.747	1.00 21.89	AAAA
	2445	N	LEU A 308	31.588	46.851	65.192	1.00 20.10	AAAA
MOTA	2446		LEU A 308	31.189	46.543	66.559	1.00 21.10	AAAA
MOTA			LEU A 308	29.671	46.340	66.644	1.00 20.99	AAAA
MOTA	2447	CB		28.897	47.656	66.674	1.00 21.54	AAAA
MOTA	2448	CG	LEU A 308		47.473	66.411	1.00 19.91	AAAA
MOTA	2449		LEU A 308	27.397			1.00 21.04	AAAA
MOTA	2450		LEU A 308	29.177	48.283	68.045		
MOTA	2451	С	LEU A 308	31.886	45.284	67.052	1.00 21.98	AAAA
ATOM	2452	0	LEU A 308	32.284	45.186	68.215	1.00 22.17	AAAA
ATOM	2453	N	ILE A 309	32.023	44.310	66.165	1.00 22:32	AAAA
ATOM	2454	CA	ILE A 309	32.658	43.069	66.544	1.00 23.12	AAAA
ATOM	2455	CB	ILE A 309	32.590	42.016	65.413	1.00 22.33	AAAA .
ATOM	2456	CG2	ILE A 309	33.356	40.787	65.827	1.00 21.76	AAAA
MOTA	2457	CG1	ILE A 309	31.140	41.678	65.061	1.00 22.16	AAAA
ATOM	2458		ILE A 309	30.366	41.037	66,166	1.00 22.01	AAAA
MOTA	2459	С	ILE A 309	34.115	43.377	66.790	1.00 24.52	AAAA
ATOM	2460	Ō	ILE A 309	34.734	42.828	67.709	1.00 25.72	AAAA
ATOM	2461	N	TRP A 310	34.673	44.253	65.957	1.00 24.70	AAAA
	2462	CA	TRP A 310	36.075	44.570	66.099	1.00 24.20	AAAA
MOTA	2463	CB	TRP A 310	36.587	45.417	64.944	1.00 23.29	AAAA
ATOM	2464	CG	TRP A 310	38.040	45.712	65.123	1.00 23.17	AAAA
MOTA		CD2		39.104	44.752	65.257	1.00 21.36	AAAA
ATOM	2465 2466		TRP A 310	40.291	45.472	65.490	1.00 20.62	AAAA
MOTA			TRP A 310	39.165	43.354	65.202	1.00 20.01	AAAA
MOTA	2467		TRP A 310	38.614	46.938	65.273	1.00 22.82	AAAA
MOTA	2468			39.967	46.803	65.497	1.00 22.30	AAAA
MOTA	2469		TRP A 310	41.521	44.845	65.668	1.00 19.91	AAAA
MOTA	2470		TRP A 310		42.734	65.381	1.00 19.08	AAAA
MOTA	2471	CZ3		40.388	43.477	65.610	1.00 19.40	AAAA
ATOM	2472		TRP A 310	41.547		67.411	1.00 25.26	AAAA
MOTA	2473	С	TRP A 310	36.318	45.279	68.109	1.00 24.71	AAAA
ATOM	2474	0	TRP A 310	37.262	44.945		1.00 26.76	AAAA
MOTA	2475	N	CYS A 311	35.467	46.247.		1.00 27.89	AAAA
MOTA	2476	CA	CYS A 311	35.608	46.975	69.007	1.00 28.98	AAAA
MOTA	.2477	CB	CYS A 311	34:548	48.081	69.113	1.00 31.89	AAAA
MOTA	2478	SG	CYS A 311	34.798	49.462	67.991	1.00 27.51	AAAA
MOTA	2479	С	CYS A 311	35.495	46.043	70.212	1.00 27.31	AAAA
MOTA	2480	0	CYS A 311	36.289	46.127	71.135	1.00 27.33	AAAA
MOTA	2481	N	GLU A 312	34.495	45.169	70.187	1.00 27.33	AAAA
MOTA	2482	CA	GLU A 312	34.246	44.210	71.250	1.00 28.55	AAAA
MOTA	2483	CB	GLU A 312	33.106	43.287	70.850	1.00 28.93	AAAA
MOTA	2484	CG	GLU A 312	31.903	43.333	71.741		
ATOM	2,485	CD	GLU A 312	32.232	42.958	73.154	1.00 29.78	AAAA
MOTA	2486	0E1		32.954	41.957	73.345	1.00 30.81	AAAA
ATOM	2487	OE2		31.754	43.653	74.071	1.00 30.79	AAAA
ATOM	2488	С	GLU A 312	35.463	43.357	71 514	1.00 28.91	AAAA
MOTA	2489	0	GLU A 312	35.822	43.110	72 662	1.00 30.57	AAAA
ATOM	2490	N	LEU A 313	36.081	42.889	70.436	1.00 29.04	AAAA
MOTA	2491	CA	LEU A 313	37.266	42.045	70.516	1.00 28.87	. AAAA
MOTA	2492	CB	LEU A 313	37.524	41.373	69.157	1.00 29.39	AAAA
ATOM	2493	CG	LEU A 313	36.548	40.311	68.644	1.00 30.32	AAAA
ATOM	2494	CDI	LEU A 313	36.910	39.872	67.215	1.00 30.26	AAAA
ATOM	2495		LEU A 313	36.582	39.114	69.593	1.00 30.42	AAAA
ATOM	2496	C	LEU A 313	38.474	42.888	70.905	1.00 27.75	AAAA
ATOM	2497	ō	LEU A 313	39.215	42.553	71.808	1.00 27.34	AAAA
ATOM	2498	N	SER A 314	38.642	43.986	70.191	1.00 27.95	AAAA
ATOM	2499	CA	SER A 314	39.736	44.927	70.376	1.00 28.62	AAAA
	2500	CB	SER A 314	39.690	45.937	69.231	1.00 27.49	AAAA
MOTA	2501	OG	SER A 314	40.703	46.904	69.343	1.00 30.12	AAAA
MOTA		C	SER A 314	39.666	45.653		1.00 29.67	AAAA
. ATOM	2502	ō	SER A 314	40.488	46.517	72.023	1.00 29.00	AAAA
ATOM	2503	N	GLY A 315	38.676	45.302	72.538	1.00 30.78	AAAA
ATOM	2504		GLY A 315	38.535	45.935	73.827		AAAA
MOTA	2505	CA	GLY A 315	38.542	47.452			AAAA
MOTA	2506	C	GLY A 315	39.142	48.091	74.647		AAAA
MOTA	2507	O	ARG A 316	37.881	48.041			AAAA.
MOTA	2508	N	THIS IS SEU	31.007	-0.041			

ATOM	2509	CA	ARG A 316	37.841	49.493	72.702	1.00 39.49	AAAA
MOTA	2510	CB	ARG A 316	38.608	49.968	71.484	1.00 39.86	AAAA
ATOM	2511	CG	ARG A 316	37.946	49.677	70.161	1.00 40.77	AAAA
MOTA	2512	CD	ARG A 316		50.226	69.077	1.00 41.47	AAAA
MOTA	2513	NE	ARG A 316		49.566	69.092	1.00 42.36	AAAA
MOTA	2514	CZ	ARG A 316		50.057	68.515	1.00 43.38	AAAA
MOTA	2515		ARG A 316		51.217	67.882	1.00 44.76	
ATOM	2516		ARG A 316		49.385	68.556	1.00 43.71	AAAA
ATOM	2517	С	ARG A 316		50.015	72.631		AAAA
ATOM	2518	õ	ARG A 316				1.00 41.54	AAAA
ATOM	2519	N	GLU A 317		49.429	71.959	1.00 42.64	AAAA
	2520	CA	GLU A 317		51.119	73.329	1.00 43.10	AAAA
MOTA	2521	CB			51.720	73.356	1.00 44.51	AAAA
MOTA	2522	CG	GLU A 317 GLU A 317		52.936	74.293	1.00 46.17	AAAA
ATOM		CD	GLU A 317		52.614	75.759	1.00 49.65	AAAA
ATOM	2523		GLU A 317		51.623	76.439	1.00 52.51	AAAA
ATOM	2524				51.251	77.607	1.00 53.37	AAAA
ATOM	2525		GLU A 317	36.444	51.214	75.831	1.00 54.14	AAAA
MOTA	2526	C	GLU A 317	34.318	52.098	71.974	1.00 43.86	AAAA
ATOM	2527	0	GLU A 317	35.067	52.532	71.108	1.00 42.46	AAAA
ATOM	2528	N	VAL A 318	33.023	51.916	71.779	1.00 44.79	AAAA
MOTA	2529	CA	VAL A 318	32.394	52.197	70.502	1.00 45.57	AAAA
ATOM	2530	CB	VAL A 318	31.098	51.369	70.324	1.00 45.36	AAAA
ATOM	2531		VAL A 318		51.558	68.924	1.00 45.44	AAAA
MOTA	2532	CG2		31.366	49.911	70.612	1.00 46.35	AAAA
MOTA	2533	С	VAL A 318	32.007	53.652	70.377	1.00 46.41	AAAA
MOTA	2534	0	VAL A 318	31.199	54.145	71.165	1.00 46.53	AAAA
MOTA	2535	N	PRO A 319	32.584	54.370	69.396	1.00 46.89	AAAA
MOTA	2536	CD	PRO A 319	33.581	54.017	68.375	1.00 46.44	AAAA
MOTA		CA	PRO A 319	32.209	55.774	69.247	1.00 47.62	AAAA
MOTA	2538	CB	PRO A 319	33.022	56.206	68.024	1.00 46.96	AAAA
MOTA	2539	CG	PRO A 319	33.161	54.922	67.251	1.00 46.38	AAAA
MOTA	2540	C	PRO A 319	30.709	55.743	68.977	1.00 48.64	AAAA
MOTA	2541	0	PRO A 319	30.236	54.860	68.262	1.00 48.61	AAAA
MOTA	2542	N	GLU A 320	29.944	56.667	69.544	1.00 49.24	AAAA
ATOM	2543	CA	GLU A 320	28.522	56.598	69.288	1.00 50.01	AAAA
ATOM	2544	CB	GLU A 320	27.720	57.330	70.363	1.00 51.15	AAAA
ATOM	2545	CG	GLU A 320	27.828	58.831	70.339	1.00 53.01	AAAA
MOTA	2546	CD	GLU A 320	26.825	59.474	71.282	1.00 54.34	AAAA
MOTA	2547		GLU A 320	25.604	59.273	71.077	1.00 54.04	AAAA
MOTA	2548		GLU A 320	27.255	60.171	72.228	1.00 55.06	AAAA
ATOM	2549	C	GLU A 320	28.206	57.168	67.921	1.00 49.78	AAAA
MOTA	2550	0	GLU A 320	27.170	56.861	67.324	1.00 49.79	AAAA
MOTA	2551	N	LÝS A 321	29.116	57.980	67.407	1.00 49.26	AAAA
ATOM	2552	CA	LYS A 321	28.906	58.589	66.109	1.00 49.20	. AAAA
MOTA	2553	CB	LYS A 321	28.873	60.106	66.251	1.00 50.38	AAAA
MOTA	> 354	CG	LYS A 321	30.234	60.674	66.634	1.00 52.88	AAAA
MOTA	555	CD	LYS A 321	30.717	60.180	68.002	1.00 53.76	AAAA
MOTA	2556	CE	LYS A 321	32.229		68.154	1.00 55.00	AAAA
MOTA	2557	NZ	LYS A 321	32.715	61.725	67.829	1.00 55.95	AAAA
ATOM	2558	Ç	LYS A 321	30.037	58.207	65.171	1.00 48.64	AAAA
MOTA	2559	Ο.	LYS A 321	31.052	57.650	65.590	1.00 48.58	AAAA
ATOM	2560	N	LEU A 322	29.854	58.511	63.894	1.00 47.78	AAAA
MOTA	2561	CA	LEU A 322	30.870	58.238	62.896	1.00 46.13	AAAA
ATOM	2562	CB	LEU A 322	30.248 ·	57.638	61.638	1.00 46.84	AAAA
MOTA	2563	CG	LEU A 322	29.240	56.504	61.848	1.00 47.71	AAAA
MOTA	2564		LEU A 322	28.788	55.998	60.491	1.00 48.02	AAAA
MOTA	25 <u>.</u> 65		LEU A 322	29.853	55.374	62.667	1.00 48.21	AAAA
ATOM	2566		LEU A 322	31.427	59.608	62.580	1.00 44.61	AAAA
MOTA	2567		LEU A 322	30.674	60.571	62.491	1.00 44.73	AAAA
ATOM	2568	N	ASN A 323	32.741	59.706	62.447	1.00 42.66	AAAA
MOTA	2569	CA	ASN A 323	33.360	60.976	62.135	1.00 41.19	AAAA
ATOM	2570		ASN A 323	34.860	60.904	62.402	1.00 41.07	AAAA
ATOM	2571		ASN A 323	35.576	60.001	61.436	1.00 41.43	AAAA
ATOM ·	2572		ASN A 323	35.117	58.901	61.147	1.00 42.46	AAAA
ATOM	2573	ND2	ASN A 323	36.720	60.449	60.943	1.00 41.77	AAAA
ATOM	2574		ASN A 323	33.068	61.223		1.00 40.76	AAAA
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### SUBSTITUTE SHEET (RULE 26)

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ATOM	2575	Ο.	ASN A	323	32.	430	60.395	60.010	1.00	40.19	AAAA
MOTA	2576	N	ASN A	324	33	523	62.352	60.129	1.00	40.11	AAAA
MOTA	2577	CA	ASN A	324	33.	268	62.699	58.735	1.00	39.99	AAAA
ATOM	2578	CB	ASN A	324	33.	711	64.128	58.472	1.00	39.54	AAAA
	2579	CG	ASN A			003	65.114	59.361	1 00	40.88	AAAA
ATOM											
ATOM	2580	ODl	ASN A	324	31.	763	65.145	59.417	1.00	40.77	AAAA
ATOM	2581	צמע	ASN A	324	33	779	65.938	60.064	1.00	40.63	AAAA
	_	-									
MOTA	2582	С	asn a	324	33.	918	61.786	57.712	1.00	40.10	AAAA
MOTA	2583	0	ASN A	324	33.	320	61.468	56.678	1.00	39.24	AAAA
										40.41	
MOTA	2584	N	LYS A			144	61.376	58.011			AAAA
MOTA	2585	CA	LYS A	325	. 35.	908	60.519	57.126	1.00	41.41	AAAA
	2586	CB	LYS A			262	60.201	57.761	1 00	42.64	AAAA
ATOM											
ATOM	258 <b>7</b>	CG	LYS A	. 325	38.	224	59.504	56.828	1.00	44.45	AAAA
ATOM	2588	CD	LYS A	325	39.	575	59.199	57.491	1.00	45.61	AAAA
										45.88	
ATOM	2589	CE	LYS A		40.		60.464	57.85 <b>0</b>			AAAA
ATOM	2590	NZ	LYS A	325	41.	717	60.151	58.404	1.00	46.27	AAAA
		C		325		124	59.248	56.856	1 00	41.56	AAAA
ATOM	2591										
ATOM	2592	0	LYS A	325		042	58.781	55.716	1.00	41.35	AAAA
MOTA	2593	N	ALA A	326	34.	524	58.703	57.906	1.00	41.32	AAAA
MOTA	2594	CA	ALA A			732	57.492	57.774		41.07	AAAA
MOTA	2595	CB	ALA A	326	33.	452	56.912	59.143	1.00	40.87	AAAA
	2596	C	ALA A	326	32	420	57.722	57.019	1 00	41.24	AAAA
MOTA											
ATOM	2597	0	ALA A	326	32.	045	56.913	56.174		40.91	aaaa
MOTA	2598	N	LYS A	327	31.	719	58.815	57.316	1.00	41.92	AAAA
						451	59.097	56.631		42.20	AAAA
ATOM	2599	CA	LYS A							•	
MOTA	2600	CB	.LYS A	. 327	29.	796	60.374	57.170	1.00	43.61	AAAA
ATOM	2601	CG	LYS A	327	29	534	60.413	58,670	1.00	45.83	AAAA
ATOM	2602	CD	LYS A	. 321		745	61.681	59.029		47.34	AAAA
MOTA	2603	CE.	LYS A	327	28.	682	61.952	60.538	1.00	48.28	AAAA
		NZ	LYS A		2 9	090	60.845	61.351	1 00	48.98	AAAA
MOTA	2604										
ATOM	2605	С	LYS A	. 327	30.	673	59.266	55.125		41.33	AAAA
ATOM	2606	0	LYS A	327	29.	879	58.797	54.309	1.00	40.78	AAAA
			GLU A			761	59.950	54.781		40.39	AAAA
MOTA	2607	N		-							
ATOM	2608	, CA	GLU A	328	32.	129	60.217	53.399		38.91	AAAA
ATOM	2609	CB	GLU A	328	33.	300	61.199	53.369	1.00	40.04	AAAA -
			GLU A				62.576	53.909		41.94	AAAA
MOTA	2610	CG				941					
ATOM	2611	CD	GLU A	328	34.	131	63.515	53.994	1.00	43.77	AAAA
ATOM	2612	OEI	GLU A	328	34.	904	63.595	53.010	1.00	44.29	AAAA
						285		55.040		45.11	AAAA
ATOM	2613	_	GLU A				64.189				
MOTA	2614	С	GLU A	. 328	32.	497	58.938	52.675	1.00	37.39	AAAA
ATOM	2615	0	GLU A	328	32.	114	58.722	51.525	1.00	37.31	AAAA
								53.355		35.67	AAAA
ATOM	2616	N	LEU A			255	58.091				
ATOM	2617	CA	LEU A	. 329	33.	657	56.820	52.783	1.00	33.03	AAAA
ATOM	2618	CB	LEU A		34	451	56.012	53.813	1.00	30.62	AAAA
										27.48	
MOTA	2619	CG	LEU A	329		760	54.549	53.481			AAAA
ATOM	2620	CD1	LEU A	329	35.	549	54.453	52.193	1.00	<b>?6.24</b>	AAAA
	2621		LEU A			514	53.936	54.622	1 00	25.74	AAAA
MOTA	-										
ATOM	2622	С	LEU A	329	32,	405	56.057	52.368	1.00	33.24	AAAA
ATOM	2623	0	LEU A	329	32.	239	55.708	51.205	1.00	32.72	AAAA
								53.327		33.92	AAAA
ATOM	2624	N	LEU A			519	55.810				
MOTA	2625	CA	LEU A	. 330	30.	289	55.090	53.046	1.00	34.91	AAAA
ATOM	2626	CB	LEU A	330	29	411	55.023	54.292	1.00	34.02	AAAA
ATOM	2627	CG	LEU A			067	54.236	55.418		34.06	AAAA
MOTA	. 2628	CD1	LEU A	. 330	29.	096	54.060	56.571	1.00	33.63	AAAA
			LEU A		3.0	512	52.892	54.884	1.00	33.82	AAAA
MOTA	2629										
MOTA	2630	С	LEU A	. 330	29.	499	55.695	51.907		35.94	AAAA
ATOM	2631	0	LEU A	. 330	28.	984	54.968	51.060	1.00	36.14	AAAA
			LYS A			415	57.022	51.883		38:17	AAAA
MOTA	2632	N									
ATOM	2633	CA	LYS A	. 331	28.	664	57.718	50.845	1.00	41.05	AAAA
ATOM	2634	CB	LYS A		- 28.		59.161	51.233	1.00	41.83	AAAA
										43.26	AAAA
MOTA	2635	CG	LYS A			584	59.358	52.497			
MOTA	2636	CD	LYS A	331	27.	202	60.823	52.755	1.00	44.15	AAAA
			LYS A			182	61.333	51,730	1.00	45.71	AAAA
ATOM	2637	CE									
MOTA	2638	NZ	LYS A			695	62.735	51.993		45.95	AAAA
MOTA	2639	С	LYS A	331	29.	342	57.681	49.490	1.00	42.22	AAAA
		ō	LYS A			712	57 000	48.480	1.00	41.94	AAAA
ATOM	2640	J	Tira ti		40.	,,2	31.300				,
		•									•

ATOM`	2641	N.	SER A 332		30.618	57.316	49,463	1.00 4	A 45	
										AAAA
MOTA	2642	CA	S <u>E</u> R A 332		31.351	57.271	48.202	1.00 4	6.88	AAAA
MOTA	2643	CB	SER A 332		32.854	57.416	48.435	1.00 4	6.49	AAAA
ATOM	2644	OG	SER A 332		33.380	56.263	49.058	1.00 4	5.65	AAAA
ATOM	2645	c	SER A 332		31.093	55.959	47.494	1.00 4		
										AAAA
- MOTA	2646	0	SER A 332		31.262	55.854	46.281	1.00 4		AAAA
ATOM	2647	N	ILE A 333		30.697	54.952	48.258	1.00 5	0.62	. AAAA
ATOM	2648	CA	ILE A 333		30.420	53.648	47.686	1.00 5	2.65	AAAA
ATOM	2649	CB	ILE A.333		30.246	52.584		1.00 5		AAAA
-										
MOTA	2650		ILE A 333		29.889	51.248	48.157	1.00 5		AAAA
MOTA	2651	CG1	ILE A 333		31.522	52.465	49.596	1.00 5	2.29	AAAA
ATOM '	2652	CD1	ILE A 333		31.403	51.463	50.696	1.00 5	3.23	AAAA
ATOM	2653	С	ILE A 333	•	29.120	53.712	46.924-	1.00 5	4.42	AAAA
ATOM	2654	ō	ILE A 333		28.122	54.178	47.462	1.00 5		AAAA
-										
ATOM	2655	N	ASP A 334		29.118	53.274	45.672	1.00 5		AAAA
ATOM	2656	CA	ASP A 334		27.863	53.263	44.940	1.00 5	9.13	- AAAA
ATOM	2657	CB	ASP A 334		28.050	53.460	43.433	1.00 5	9.64	AAAA
ATOM	2658	CG	ASP A 334		28.976	52.446	42.823	1.00 5	9.23	AAAA
ATOM	2659		ASP A 334		28.853	52:194	41.606	1.00 5		· AAAA
							43.559			
ATOM	2660		ASP A 334		29.839	51.925		1.00 5		AAAA
ATOM	2661	Ç	ASP A 334		27.251	51.898	45.215	1.00 6		AAAA
MOTA	2662	0	ASP A 334		27.803	50.861	44.840	1.00 6	1.15	AAAA
MOTA	2663	N	PHE A 335		26.113	51.914	45.897	1.00 6	2.56	AAAA
MOTA	2664	CA	PHE A 335		25.414	50.701	46.257	1.00 6		AAAA
			PHE A 335		25.311	50.621	47.779	1.00 6		
ATOM .	2665	CB								AAAA
MOTA	2666	ÇG	PHE A 335		24.224	49.714	48.263	1.00 6		AAAA
MOTA	2667	CD1	PHE A 335		24.180	48.379	47.868	1.00 6	5.54	AAAA
ATOM	2668	CD2	PHE A 335		23.234	50.197	49.107	1.00 6	5.12	AAAA
MOTA	2669	CE1	PHE A 335		23.163	47.539	48.305	1.00 6	5.75	AAAA
ATOM	2670		PHE A 335		22.213	49.367	49.552	1.00 6		AAAA
	2671	CZ	PHE A 335	•	22.177	48.034	49.150	1.00 6		AAAA
MOTA							45.640	-		
MOTA	2672	Ċ	PHE A 335		24.025	50.626		1.00 6		AAAA
ATOM	2673	0	PHE A 335		23.591	49.564	45.184	1.00 6		AAAA
MOTA	2674	N	GLU A 336		23.338	51.763	45.618	1.00 6	6.38	AAAA
MOTA	2675	CA	GLU A 336		21.980	51.826	45.097	1.00 6	7.49	AAAA
MOTA	2676	CB	GLU A 336		21.893	51.26Ò	43.673	1.00 6	8.25	AAAA
ATOM	2677	CG	GLU A 336		20.459	51.230	43.116	1.00 6	9.15	AAAA
ATOM	2678	CD	GLU A 336		20.334	50.465	41.804	1.00 6		AAAA
			GLU A 336		20.710	49.271	41.784	1.00 6		AAAA
ATOM	2679									
Mota	2680		GLU A 336		19.851	51.051	40.804	1.00 6		AAAA
MOTA	2681	С	GLU A 336		21.098	50.999	46.025	1.00 6		AAAA
MOTA	2682	0	GLU A 336		21.216	49.776	46.082	1.00 6	7.58	AAAA
ATOM	2683	N	GLU A 337		20.227	51.679	46.761	1.00 6	7.87	AAAA
ATOM	2684	CA	GLU A 337		19.317	51.020	47.686	1.00 6	8.66	AAAA
	2685	CB	GLU A 337		18.583	52.085	48.502	1.00 6		AAAA
ATOM						51.715				AAAA
ATOM	2686	CG.	GLU : 337		18.279		49.944			
ATOM	2687	CD	GLU. A 337		19.527	51.587	50.789	1.00 6		AAAA
ATOM	2688	OE1	GLU 337		20.319	52.554	50.851	1.00 6	7.05	AAAA
ATOM	2689	OE2	GLU A 337		19.711	50.518	51.398	1.00 6	7.79	AAAA
ATOM	2690	С	GLU A 337		18.322	50.222	46.827	1.00 6	9.28	AAAA
ATOM	2691	ō	GLU A 337		17.886	50.705	45.780	1.00 6		AAAA
							47.259	1.00 6		AAAA
MOTA	2692	N	PHE A 338		17.966	49.012				
MOTA	2693	CA	PHE A 338		17.035	48.176	46.497	1.00 6		ААДА
ATOM	2694	CB	PHE A 338		16.995	46.759	47.066	1.00 7		AAA'A
ATOM	2695	CG	PHE A 338		16.225	45.789	46.221	1.00 7	1.57	AAAA
ATOM	2696	CD1	PHE A 338		16.666	45.462	44.936	1.00 7	2.04	AAAA
MOTA	2697		PHE A 338		15.052	45.208	46.698	1.00 7		AAAA
					15.944		44.138	1.00 7		AAAA
MOTA	2698		PHE A 338			44.566				
ATOM	2699		PHE A 338		14.323	44.313	45.909	1.00 7		AAAA
MOTA	2700	CZ	PHE A 338		14.770	43.991	44.627	1.00 7		AAAA
ATOM	2701	С	PHE A 338		15.633	48.770	46.494	1.00 6	9.26	AAAA
ATOM	2702	ō	PHE A 338		15.072	49.029	45.434	1.00 6		AAAA
	2703	N	ASP A 339		15.053	48.962	47.674	1.00 6		AAAA
MOTA			ASP A 339		13.733		47.755	1.00 6		· AAAA
MOTA	2704	CA				49.572				
ATOM	2705	CB	ASP A 339		13.134	49.457	49.157	1.00 6		AAAA
MOTA	2706	CG	ASP A 339		11.819	50.233	49.299	1.00 6	9.72	AAAA

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### **257/263** Figure 19-42

WO 01/018045

							1 00 50 30	2222
MOTA	2707	OD1	ASP A 339	11.813	51.462	49.058	1.00 69.39	AAAA
-	2708	OD2	ASP A 339	10.790	49.618	49.655	1.00 69.78	AAAA
MOTA			ASP A 339	13.972	51.035	47.440	1.00 69.95	AAAA
MOTA	2709	С				48.333	1.00 69.92	AAAA
MOTA	2710	0	ASP A 339	14.305	51.815			
ATOM	2711	N	ASP A 340	13.810	51.389	46.168	1.00 70.23	AAAA
	2712	CA	ASP A 340	14.023	52.748	45.699	1.00 70.39	AAAA
MOTA				12.757	53.283	45.041	1.00 70.64	AAAA
ATOM	2713	CB	ASP A 340			43.791	1.00 70.86	AAAA
ATOM	2714	CG	ASP A 340	12.397	52.517			
MOTA	2715	0D1	ASP A 340	12.126	51.302	43.903	1.00 70.93	AAAA
		002	ASP A 340	12.399	53.125	42.699	1.00 70.89	AAAA
MOTA	2716		722 7 740		53.674	46.807	1.00 70.63	AAAA
ATOM	2717	С	ASP A 340	14.482			1.00 71.13	AAAA
ATOM	2718	0	ASP A 340	15.688	53.847	47.008		
	2719	N	GLU A 341	13.543	54.259	47.544	1.00 69.95	AAAA ·
ATOM	-		GLU A 341	13.947	55.150	48.619	1.00 69.17	AAAA
MOTA	2720	CA			56.613	48,266	1.00 70.83	AAAA
ATOM	2721	CB	GLU A 341	13.636			1.00 73.44	AAAA
MOTA	2722	CG	GLU A 341	14.098	57.601	49.347		
	2723	CD	GLU A 341	13.956	59.071	48.951	1.00 75.27	AAAA
ATOM			GLU A 341	12.825	59.518	48.646	1.00 76.21	AAAA
ATOM	2724					48.954	1.00 75.69	AAAA
ATOM	2725	OE2	GLU A 341	14.984	59.786			AAAA
MOTA	2726	С	GLU A 341	13.367	54.819	49.983	1.00 67.09	
	2727	ō	GLU A 341	12.233	55.176	50.297	1.00 66.57	AAAA
ATOM				14.158	54.114.	50.785	1.00 64.87	AAAA
MOTA	2728	N	VAL A 342			52.148	1.00 62.55	AAAA
ATOM	2729	CA	VAL A 342	13.767	53.779			AAAA
ATOM	2730	CB	VAL A 342	14.265	52.377	52.589	1.00 62.81	
	2731		VAL A 342	14.042	52.193	54.081	1.00 62.56	AAAA
MOTA			VAL A 342	13.513	51.298	51.849	1.00 63.69	AAAA
MOTA	2732					52.982	1.00 59.94	AAAA
ATOM	2733	С	VAL A 342	14.483	54.822			AAAA
ATOM	2734	0	VAL A 342	14.022	55.215	54.054	1.00 59.91	
MOTA	2735	N	ASP A 343	15.609	55.278	52.442	1.00 56.85	AAAA
			ASP A 343	16.457	56.266	53.085	1.00 54.01	AAAA
MOTA	2736	CA			57.446	53.605	1.00 54.18	AAAA
ATOM	2737	CB	ASP A 343	15.639			1,00 53.96	AAAA
ATOM	2738	CG	ASP A 343	16.505	58.511	54.241		
ATOM	2739	OD1	ASP A 343	15.947	59.485	54.785	1.00 54.59	AAAA
			ASP A 343	17.747	58.373	54.191	1.00 53.61	AAAA
MOTA	2740			17.186	55.609	54.242	1.00 51.92	AAAA
MOTA	2741	С	ASP A 343			55.307	1.00 51.89	AAAA
ATOM	2742	0	ASP A 343	16.611	55.371			AAAA
MOTA	2743	N	ARG A 344	18.458	55.306	54.029	1.00 48.86	
	2744	CA	ARG A 344	19.240	54.676	55.069	1.00 45.59	AAAA
MOTA				19.847	53.369	54.573	1.00 43.94	AAAA
MOTA	2745	CB	ARG A 344			54.220	1.00 41.70	AAAA
MOTA	2746	CG	ARG A 344	18.847	52.289		1.00 38.94	AAAA
MOTA	2747	CD	ARG A 344	17.953	51.955	55.385	1.00 36.54	
	2748	NE	ARG A 344	17.139	50.781	55.096	1.00 36.78	AAAA
ATOM			ARG A 344	16.176	50.316	55.888	1.00 34.81	AAAA
ATOM	2749	CZ			50.927	57.033	1.00 34.11	AAAA
MOTA	2750	NH1	. ARG A 344	15.890			1.00 31.84	·AAAA
ATOM	2751	NH2	ARG A 344	15.506				
	2752	С	ARG A 344	20.340	55.604	55.520	1.00 44.83	AAAA
ATOM	2753	ō	ARG A 344	21.308	55.157	56.128	1.00 43.97	A AA
MOTA				20.192			1.00 44.32	A. AA
ATOM	2754	N	SER A 345				1.00 43.74	AAAA
MOTA	2755	CA	SER A 345	21.199				AAAA
ATOM	2756	CB	SER A 345	20.860	59.248		1.00 44.49	
	2757		SER A 345	19.645	59.729	55.577	1.00 46.07	AAAA
ATOM				21.307	57.977	57.144	1.00 42.82	AAAA
MOTA	2758	С	SER A 345				1.00 42.91	AAAA
ATOM	2759	0	SER A 345	22.304			1.00 41.40	
ATOM	2760		TYR A 346	20.282	57.509	57.849	1.00 41.48	AAAA
			TYR A 346	20.296	57.549	59.303	1.00 40.35	AAAA
MOTA	2761			18.947			1.00 40.38	AAAA
ATOM	2762	CB	TYR A 346				1.00 39.28	AAAA
MOTA	2763	CG	TYR A 346	18.630			1.00 33.40	
	2764	CD1	TYR A 346	19.293	54.589			AAAA
MOTA			TYR A 346	19.022		60.079	1.00 37.71	AAAA
ATOM	2765							AAAA
MOTA	2766			17.682				AAAA
ATCM	2767	CE		17.405				
	2768		TYR A 346	18.079	52.899	59.126		AAAA
ATOM			TYR A 346	17.794			1.00 37.14	AAAA
MOTA	2769							AAAA
ATOM	2770	) C	TYR A 346	21.436				AAAA
MOTA	2771		TYR A 346	21.967				
	2772		MET A 347	21.800	55.640	59.113	1.00 39.14	AAAA
ATOM	2112					•		•
•								

### SUBSTITUTE SHEET (RULE 26)

	2772	~ .	MEG		242	22.879	54.756	59.530	1.00 38.19	***
MOTA	2773	CA	MET							AAAA
MOTA	2774	CB	MET	Α	347	23.042	53.582	58.566	1.00 38.26	AAAA
MOTA	2775	CG	MET	Α	347	21.973	52.523	58.694	1.00 38.17	AAAA
	2776	SD			347	22.317	51.115	57.641	1.00 38.05	AAAA
ATOM									•	
MOTA	2777	CE			347	22.237	51.892	56.101	1.00 37.61	AAAA
ATOM	2778	C	MET	A	347	24.189	55.494	59.603	1.00 38.00	AAAA
ATOM	2779	0	MET	A	347	25.127	55.033	60.250	1.00 37.40	AAAA
					348	24.248		58.929	1.00 38.08	AAAA
ATOM	2780	N					56.637			
ATOM	2781	CA	LEU	Α	348	25.449	57.463	58.898	1.00 38.07	AAAA
MOTA	2782	CB	LEU	Α	348	25.445	58.330	57.638	1.00 36.66	AAAA
ATOM	2783	CG	LEU	Α	348	25.379	57.583	56.310	1.00 35.47	AAAA
			LEU					55.165	1.00 34.51	
ATOM	2784					25.285	58.559			AAAA
ATOM	2785	CD2	LEU	A	348	26.605	56.716	56.167	1.00 36.56 .	AAAA
ATOM-	2786	С	LEU	Α	348	25.521	58.353	60.138	1.00 39.07	AAAA
ATOM	2787	0	LEH	Α	348	26.546	58.980	60.406	1.00 38.81	AAAA
					349				1.00 39.90	
MOTA	2788	И.				24.432	58.385	60.898		AAAA
ATOM	2789	CA	GLU	Α	349	24.363	59,213	62.092	1.00 40.95	· AAAA
MOTA	2790	CB	GLU	Α	349	22.961	59.821	62.203	1.00 41.70	AAAA
ATOM	2791	CG			349	22.515	60.629	60.966	1.00 42.28	AAAA
									1.00 42.51	
ATOM	2792	CD			349	23.349	61.891	60.708		AAAA
ATOM	2793		GLU			23.414	62.778	61.587	1.00 42.38	AAAA
ATOM	2794	OE2	GLU	A	349	23.933	61.998	59.614	1.00 43.34	AAA:A
ATOM	2795	C			349	24.740	58.511	63.406	1.00 41.12	AAAA
			- :						1.00 41.38	AAAA
ATOM	2796	0			349	24.664	59.118	64.476	the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the s	
ATOM	2797	N	THR	A	350	25.140	57.243	63.326	1.00 40.86	AAAA
ATOM	2798	CA	THR	A	350	25.555	56.475	64.504	1.00 40.69	AAAA
ATOM	2799	CB	THR	A	350	24.405	56.283	65.510	1.00 41.56	AAAA
			THR			24.062	57.549	66.078	1.00 41.48	AAAA
ATOM	2800									
MOTA	2801	CG2	THR			24.821	55.345	66.638	1.00 41.19	AAAA
ATOM	2802	С	THR	Α	350 ,	26.109	55.109	64.141	1.00 40.14	AAAA
ATOM	2803	0	THR	Α	350	25.857	54.595	63.055	1.00 39.93	AAAA
					351	26.865	54.527	65.067	1.00 40.32	AAAA
MOTA	2804	N								
MOTA	2805	CA			351	27.491	53.227	64.857	1.00 40.70	AAAA
ATOM	2806	CB	LEU	Α	351	28.855	53.213	65.540	1.00 39.89	AAAA
ATOM	2807	CG	LEU	А	351	29.911	52.290	64.951	1.00 39.68	AAAA
ATOM	2808		LEU			31.170	52.403	65.772	1.00 39.88	AAAA
2.4						29.414	50.861	64.945	1.00 40.70	AAAA
MOTA	2809		LEU							
MOTA	2810	С			351	26.612	52.091	65.384	1.00 41.12	AAAA
MOTA	2811	0	LEU	A	351	26.467	51.060	64.736	1.00 40.02	AAAA
ATOM	2812	N·	LYS	A	352	26.040	52.292	66.567	1.00 42.99	AAAA
					352	25.138	51.326	67.201	1.00 43.93	AAAA
MOTA	2813	CA								
ATCM	2814	CB			352	25.412	51.225	68.707	1.00 43.38	AAAA
ATOM	2815	CG	LYS	Α	352	26.743	50.597	69.055	1.00 44.68	AAAA
MOTA	2816	CD	LYS	Α	352	27.185	50.927	70.482	1.00 45.48	AAAA
	2817	CE			352	26.189	50.500	71.539	1.00 46.21	AAAA
MOTA						36.646		72.895	1.00 47.34	AAAA
ATOM	2818	NZ	LYS				50.944		•	
ATOM	2819	C			352	23.723	51.838	67.003	1.00 44.40	AAAA
ATCM	2820	0	LYS	A	352	23.375	52.917	67.488	1.00 45.79	AAAA
ATOM	2821	N	ASP	Α	353	22.904	51.083	66.287	1.00 44.78	AAAA
	2822					21.532	51.509	66.074	1.00 44.79	AAAA
MOTA		CA			353					
MOTA	2823	CB	ASP			21.050	51.030	64.702	1.00 45.20	AAAA
ATOM	2824	CG	ASP	A	353	21.146	49.544	64.546	.1.00 45.21	AAAA
ATOM	2825		ASP			21.806	49.086	63.581	1.00 45.06	AAAA
		002	ASP	n	353	20.549	48.841	65.391	1.00 45.54	AAAA
MOTA	2826		ASP	^	333					
MOTA	2827	C	ASP			20.645	50.993	67.217	1.00 44.44	AAAA
ATCM	2828	0	ASP	A	353	21.042	50.113	67.973	1.00 44.29	AAAA
ATOM	2829	N	PRO			19.439	51.553	67.367	1.00-44.22	AAAA
	2830		PRO			18.839	52.617	66.550	1.00 44.38	AAAA
ATOM		CD								AAAA
ATOM	2831	CA	PRO			18.500	51.163	68.419	1.00 44.18	
ATCM	2832	CB	PRO			17.371	52.170	68.238	1.00 44.52	AAAA
ATCM	2833	CG	PRO			17.368	52.341	66.749	1.00 44.66	AAAA
			PRO	2	354	17.995	49.740	68.328	1.00 43.87	AAAA
ATCM	2834	C	FRU		254				1.00 44.48	AAAA
ATOM .	2835	0	PRO			17.962	49.152	67.249		
ATCM -	2836	N	TRP			17.588	49.198	69.469	1.00 43.23	AAAA
ATOM	2837	CA	TRP			17.051	47.851	69.500	1.00 42.88	AAAA
		CB	TRP			16.743	47.401	70.927	1.00 46.42	AAAA.
ATCM	2838	CB	TKP	••		20.730	47.4			•

ATOM	2839	CG	TRP A 35	17.959	47.052	71.695	1.00 49.91	AAAA
MOTA	2840	CD2	TRP A 35!	18.476	45.733	71.903	1.00 51.56	AAAA
ATOM	2841	CE2	TRP A 35!	19.684	45.868	72.627	1.00 52.03	AAAA
MOTA	2842	CE3	TRP A 35!	18.038	44.450	71.548	1.00 52.25	AAAA
ATOM	2843	CD1	TRP A 355	18.846	47.915	72.284	1.00 50.53	AAAA
ATOM	2844	NE1	TRP A 35!	19.885	47208	72.846	1.00 51.63	AAAA
MOTA	2845	CZ2	TRP A 35!	20.460	44.763	73.003	1.00 52.64	AAAA
MOTA '	2846	CZ3	TRP A 35	18.810	43.352	71.921	1.00 53.12	AAAA
ATOM	2847	CH2	TRP A 35!	20.008	43.518	72.642	1.00 53.02	AAAA
						68.675	1.00 40.28	
ATOM	2848	C	TRP A 35		47.767			AAAA
ATOM	2849	0	TRP A 35	15.017	48.720	68.591	1.00 39.82	AAAA
ATOM	2850	N	ARG A 35	15.591	46.610	68.065	1.00 36.83	AAAA
MOTA	2851	CA	ARG A 35	14.440	46.365	67.225	1.00 33.70	AAAA
MOTA	2852	ÇВ	ARG A 35	14.901	46.197	65.772	1.00 29.50	AAAA
N COM	2853	CG	ARG A 35	15.635	47.423	65.256	1.00 25.22	AAAA
MOTA	-							
ATOM	2854	CD	ARG A 35	16.418	47.194	63.973	1.00 21.53	AAAA
MOTA	2855	NE	ARG A 35	17.055	48.435	63.533	1.00 18.55	AAAA
MOTA	2856	CZ	ARG A 35	17.976	48.533	62.574	1.00 17.06	AAAA
MOTA	2857	NHl	ARG A 35	18.403	47.451	61.919	1.00 17.64	AAAA
MOTA	2858	NH2	ARG A 35	18.445	49.721	62.241	1.00 11.56	AAAA
MOTA	2859	С	ARG A 35	13.831	45.095	67.773	1.00 34.63	AAAA
	2860		ARG A 35	13.605	44.117	67.051	1.00 35.86	AAAA
ATOM		0						
ATOM	2861	N	GLY A 35'	13.587	45.112	69.079	1.00 34.58	AAAA
						69.734		
ATOM	2862	CA	GLY A 35		43.960		1.00 34.33	AAAA
ATOM	2863	С	<b>GLY A 35</b>	11.536	43.783	69.395	1.00 34.31	AAAA
ATOM	2864	0	<b>GLY A 35</b> °	11.006	44.418	68.484	1.00 33.56	AAAA
MOTA	2865	N	GLY A 35	10.876	42.906	70.139	1.00 34.47	AAAA
MOTA	2866	CA	GLY A 35	9.468	42.656	69.916	1.00 34.61	AAAA
MOTA	2867	С	GLY A 35	9.114	41.389	70.655	1.00 34.47	AAAA
ATOM			•					
ATOM -	2868	0	GLY A 35	9.962	40.821	71.345	1.00 34.27	AAAA
	2060	N	GLU A 35	7.869	40.948	70.523	1.00 34.16	AAAA
MOTA	2869		-					
ATOM	2870	CA	GLU A 35:	7.438	39.729	71.180	1.00 33.94	AAAA
			GLU A 35		39.644	71.174	1.00 34.78	AAAA
ATOM	2871	CB						
MOTA	2872	ÇG	GLU A 35:	5.278	40.648	72.123	1.00 36.70	AAAA
								AAAA
MOTA	2873	CD	GLU A 35	3.863	41.020	71.740	1.00 38.40	
MOTA	2874	OEL	GLU A 35	3.017	40.108	71.600	1.00 39.65	AAAA
MOTA	2875	OE2	GLU A 35	3.598	42.234	71.584	1.00 38.52	AAAA
ATOM	2876	С	GLU A 35	8.058	38.549	70.464	1.00 32.86	AAAA
ATOM	2877	0	<b>GLU A 35</b> :	8.678	38.692	69.427	1.00 32.92	AAAA
	2878	N	VAL A 36	7.918	37.375	71.036	1.00 32.63	AAAA
ATOM								
ATOM	2879	CA	VAL A 36	8.480	36.215	70.409	1.00 32.70	AAAA
	2880	CB	VAL A 36	9.422	35.472	71.376	1.00 33.24	AAAA
MOTA								
ATOM	2881	CG1	VAL A 36	10.017	34.252	70.701	1.00 32.99	AAAA
	2882	CG2	VAL A 36	10.521	36.406	71.827	1.00 32.09	AAAA
MOTA								
MOTA	2883	С	VAL A 36	7.339	35.319	69.976	1.00 32.81	AAAA
	2884	0	VAL A 36	6.702	34.660	70.791	1.00 32.02	· AAAA
MOTA								
ATOM	2885	N	ARG A 36.	7.084	35.321	68.674	1.00 33.12	AAAA
		CA	ARG A 36		34.508	68.086	1.00 33.52	AAAA
MOTA	2886							
MOTA	2887	CB	ARG A 36.	6.148	34.558	66, 565	1.00 33.43	AAAA
		CG	ARG A 36		35.885	65.967	1.00 34.35	AAAA
MOTA	2888							
MOTA	2889	CD	ARG A 36	6.041	35.972	64.469	1.00 33.90	AAAA
						64.193	1.00 31.70	AAAA
ATOM	2890	NE	ARG A 36	7.430	36.331		1.00 31.70	
ATOM	2891	CZ	ARG A 36	7.890	36.608	62.978	1.00 31.18	AAAA
ATOM	2892		ARG A 36.		36.562	61.941	1.00 30.48	AAAA
	2893	NHO	ARG A 36	9.162	36.948	62.802	1.00 29.71	AAAA
MOTA								
ATOM	2894	С	ARG A 36	6,066	33.057	68.557	1.00 34.20	AAAA
		0	ARG A 36		32.537	68.968	1.00 33.79	AAAA
MOTA	2895							
ATOM	2896	N	LYS A 36	4.914	32.407	68.496	1.00 34.68	AAAA
					31.022	68.901	1.00 35.62	AAAA
ATOM	2897	CA	LYS A 36					
	2898	CB	LYS A 36	3.350	30.555	68.782	1.00 37.80	AAAA
ATOM								
ATOM	2899	CG	LYS A 36		31.226	69.756	1.00 40.38	AAAA
	2900	CD	LYS A 36		32.777	69.77 <b>7</b>	1.00 42.09	AAAA
ATOM								
ATOM	2901	CE	LYS A 36	2.208	33.446	68.420	1.00 41.94	AAAA
		NZ	LYS A 36		34.909	68.451	1.00 39.85	AAAA
ATOM	2902						_	
ATCM	2903	С	LYS A 36	5.710	30.177	68.005	1.00 35.12	AAAA
		0.	LYS A 36		29.301	68.487	1.00 34.14	AAAA
ATOM	2904	Ф.		· v.azj	20.00			

3 mov	2905	N .	GLU A	363	5.661	30.460	66.703	1.00 35.12	AAAA
MOTA MOTA	2905	CA	GLU A		6.445	29.741	65.699	1.00 35.62	AAAA
	2907	CB	GLU A		6.567	30.560	64.424	1.00 36.81	AAAA
ATOM		CG	GLU A		5.280	30.808	63.711	1.00 38.66	AAAA
ATOM	2908	CD	GLU A		5.477	31.704	62.517	1.00 39.60	AAAA
ATOM	2909		GLU A		6.287	31.324	61.637	1.00 39.11	AAAA
ATOM -	2910		GLU A		4.826	32.782	62.469	1.00 39.95	AAAA
ATOM	2911		GLU A		7.836	29.450	66.181	1.00 35.14	ĀAAA
ATOM	2912	C	GLU A		8.321	28.316	66.098	1.00 34.50	AAAA
MOŢA	2913	0			8.475	30.505	66.671	1.00 34.96	AAAA
ATOM	2914	N	VAL A		9.830	30.431	67.180	1.00 34.44	AAAA
MOTA	2915	CA	VAL A		10.338	31.821	67.570	1.00 33.68	AAAA
ATOM	2916	CB	VAL A	•	10.338	31.722		1.00 34.13	AAAA
MOTA	2917		VAL A				66.347	1.00 31.83	AAAA
MOTA	2918		VAL A		10.337	32.715		1.00 34.44	AAAA
ATOM	2919	C	VAL A		9.908	29.499	68.370	1.00 36.01	-AAAA
MOTA	2920	0	VAL A		10.789	28.640	68.430	1.00 33.27	AAAA
ATOM	2921	N	LYS A		8.980	29.649	69.305	1.00 33.20	AAAA
MOTA	2922	CA	LYS A		8.970	28.790	70.476		AAAA
ATOM	2923	CB	LYS A		7.968	29.319	71.508	1.00 34.28	
MOTA	2924	CG	LYS A		8.307	30.705	72.033	1.00 33.67	AAAA
ATOM	2925	CD	LYS A		7.282	31.181		1.00 34.85	AAAA
ATOM	2926	CE	LYS A		7.658	32.534	73.638	1.00 36.47	AAAA
ATOM	2927	NZ	LYS A		6.698	32.990	74.710	1.00 37.91	AAAA
MOTA	2928	С	LYS A		8.654	27.342	70.109	1.00 32.75	AAAA
ATOM	2929	0	LYS A		9.071	26.421	70.818	1.00 31.95	AAAA
MOTA	2930	N	ASP A		7.919	27.136	69.012	1.00 32.81	AAAA
MOTA	2931	CA	ASP A		7.600	25.777	68.581	1.00 33.65	AAAA
MOTA	2932	CB	ASP A		6.459	25.726	67.557	1.00 33.98	AAAA
ATOM	2933	CG	ASP A		5.131	26.107	68.140	1.00 33.94	AAAA
MOTA	2934		ASP A		4.870	25.767	69.307	1.00 33.89	AAAA
MOTA	2935		ASP A		4.332	26.722	67.412	1.00 35.08	AAAA
ATOM	2936	С	ASP A		8.820	25.167	67.940	1.00 33.05	AAAA
ATOM	2937	0	ASP A		9.140	24.006	68.172	1.00 33.66	AAAA
ATOM	2938	N	THR A		9.473	25.959	67.102	1.00 33.07	AAAA
ATOM	2939	CA	THR A		10.684	25.540	66.412	1.00 32.27	AAAA
ATOM	2940	CB	THR A		11.304	26.719	65.641	1.00 32.28	AAAA
ATOM	2941		THR A		10.473	27.039	64.520	1.00 30.64	AAAA
ATOM	2942		THR A		12.711	26.377	65.166	1.00 33.29	AAAA AAAA
MOTA	2943.	C	THR A		11.680	25.044	67.442	1.00 31.71	AAAA
MOTA .	2944	0	THR A		12.178	23.918	67.352	1.00 30.45	
ATOM	2945	N	LEU A		11.955	25.896	68.426	1.00 32.05	AAAA
ATOM	2946	CA	LEU A		12.888	25.560	69.482	1.00 32.49	AAAA
ATOM	2947	CB		368		26.749	70.421	1.00 32.27	AAAA AAAA
ATOM	2948	CG	LEU A		14.097	27.809	69.960	1.00 32.71	AAAA
ATOM	2949		LEU A		15.488	27.170	69.899	1.00 33.00	AAAA
ATOM	2,950		LEU A		13.709	28 393	68.597	1.00 31.86	
MOTA	2951	С	LEU A		12.455	24.334	70.256	1.00 33.82	
ATOM	2952	0	LEU A		13.266	23.133	70.489	1.00 34.29	
MOTA	2953	N	GLU A		11.183	24.285	70.645	1.00 34.30	
ATOM	2954	CA	GLU A		10.687	23.135	71.375	1.00 36.07	
ATOM	2955	CB	GLU A		9.211	23.319	71.748	1.00 38.71	
ATOM	2956	CG	GLU A		8.974	24.285	72.920	1.00 40.18	AAAA
ATOM	2957	CD	GLU A		7.509	24.359	73.341	1.00 41.56	
ATOM	.2958		GLU A		6.917	23.276	73.572	1.00 41.83	AAAA
ATOM	2959	OE2	GLU A		6.957	25.489	73.451	1.00 41.74	AAAA
MOTA	2960	С	GLU A		10.893	21.822	70.611	1.00 37.16	AAAA
ATOM	2961	0	GLU A		11.338	20.831	71.196	1.00 37.00	AAAA
ATOM	2962	N	LYS A		10.586	21.788	69.315	1.00 37.71	AAAA
ATOM	2963	CA	LYS A		10.797	20.547	68.5 <b>67</b>	1.00 38.46	AAAA
MOTA	2964	CB	LYS A	370	10.166	20.604	67.177	1.00 39.96	AAAA
ATOM	2965	CG	LYS A	370	8.646	20.532	67.186	1.00 42.68	AAAA
ATOM	2966	CD	LYS A		8.092	20.320	65.775	1.00 44.54	AAAA
ATOM	2967	CE	LYS A	370	6.572	20.075	65.781	1.00 45.55	
MOTA	2968	NZ	LYS A	370	6.009	19.797	64.409	1.00 45.50	
ATOM	2969	C	LYS A	370	12.282	20.235	68.452	1.00 38.34	
MOTA	2970	0	LYS A	370	12.683	19.071	68.493	1.00 37.86	
							_		•

ATOM	2971	N	ALA	Α	371	13.105	21.266	68.311	1.00 37.69	AAAA
ATOM	2972	CA	ALA			14.543	21.057	68.226	1.00 37.20	AAAA
	2973	CB	ALA			15.258	22.375	67.936	1.00 35.48	AAAA
ATOM										
MOTA	2974	C			371	15.023	20.477	69.558	1.00 37.63	AAAA
MOTA	2975	0	ALA	A	371	15.920	19.626	69.585	1.00 37.12	AAAA
MOTA	2976	N	LYS	A	372	14.426	20.930	70.665	1.00 37.59	AAAA
ATOM	2977	CA	LYS			14.796	20.432	.71.995	1.00 37.46	AAAA
	2978		·LYS			14.022	21.156	73.095	1.00 36.52	AAAA
MOTA										
MOTA	2979	CG	LYS			14.287	22.634	73.111	1.00 22.67	AAAA
MOTA	2980	CD	LYS	Α	372	13.309	23.396	74.022	1.00 22.67	AAAA
MOTA	2981	CE	LYS	Α	372	13.600	24.874	73.901	1.00 22.67	AAAA
ATOM	2982	NZ	LYS	Α	372	12.692	25.708	74.785	1.00 22.67	AAAA
MOTA	2983	C	LYS			14.495	18.957	72.077	1.00 37.60	AAAA
									1.00 37.26	
ATOM	2984	0			372	15.367	18.171	72.407		AAAA
MOTA	2985	N	ALA			13.249	18.595	71.789	1.00 38.17	AAAA
MOTA	2986	CA	ALA	Α	373	12.812	17.206	71.829	1.00 39.55	AAAA
ATOM	2987	CB	ALA	Α	373	11.365	17.109	71.395	1.00 39.34	AAAA
ATOM	2988	С	AT.A	Α	373	13.675	16.277	70.972	1.00 41.02	AAAA
	2989	ŏ			373	14.366	15.410	71.561	1.00 42.26	AAAA
MOTA		_							1.00 42.06	
MOTA	2990		ALA			13.663	16.416	69.725		AAAA
HETATM	2991	ZN	ZN		951	23.696	34.788	54.072	1.00 27.38	ZONE
HETATM	2992	01	SHA	C	1	24.578	33.295	53.458	1.00 31.95	SAHA
HETATM	2993	02	SHA	С	1	24.294	35:218	51.444	1.00 33.51	SAHA
HETATM		N1	SHA	C	1	24.578	33.085	52.069	1.00 34.03	SAHA
HETATM		CI	SHA		1	24.063	34.053	51.246	1.00 34.25	SAHA
								50.259	1.00 36.87	
HETATM		C2	SHA		1	23.090	33.625			SAHA.
HETATM		C3	SHA		1	23.548	33.781	48.816	1.00 39.33	SAHA
HETATM	2998	C4	SHA	С	1	22.498	33.274	47.852	1.00 40.86	SAHA
HETATM	2999	C5	SHA	С	1	21.590	34.413	47.455	1.00 43.37	SAHA
HETATM	3000	C6	SHA	С	1	21.061	34.017	46.092	1.00 46.72	SAHA
HETATM		C7	SHA		1	19.754	34.714	45.787	1.00 48.75	SAHA
		C8	SHA		ī	19.960	35.720	44.693	1.00 50.75	SAHA
HETATM							35.467		1.00 51.08	
HETATM		03	SHA		1	20.381		43.575		SAHA
HETATM	3004	N2	SHA		1	19.591	36.956	45.085	1.00 52.52	SAHA
HETATM	3005	C9	SHA	С	1	19.842	38.330	44.507	1.00 54.25	SAHA
HETATM	3006	C10	SHA	С	1	19.243	39.431	45.215	1.00 55.76	SAHA
HETATM		C11	SHA	С	1	19.423	40.804	44.727	1.00 56.53	Saha
HETATM		C12	SHA		1	20.169	41.085	43.545	1.00 56.58	SAHA
			SHA		ī	20.755	39.942	42.827	1.00 55.93	SAHA
HETATM										
HETATM			SHA		1	20.612	38.546	43.304	1.00 54.65	SAHA
HETATM			WAT		2	36.485	44.023	49.378	1.00 4.67	SOLV
HETATM	3012	OH2	WAT	D	3	27.702	16.865	62.162	1.00 4.67	SOLV
HETATM	3013	OH2	TAW	D	4	23.251	30.387	59.575	1.00 10.12	SOLV
HETATM		OH2	WAT	D	5	33.825	41.862	46.926	1.00 21.13	SOLV
HETATM			WAT		6	24.866	44.453	47.867	1.00 23.72	SOLV
H. TATM			WAT		7	34.145	20.442	33.590	1.00 18.19	SOLV
						7.921			1.00 20.79	SOLV
HETATM		_	WAT		8		29.753	62.099		
HL-TATM			WAT		9	17.863	6.978	64.018	1.00 28.94	SOLV
HETATM	3019	OH2	WAT	D	10	35.580	44.610	74.823	1.00 31.62	SOLV
HETATM	3020	OH2	TAW	D	11	49.208	27.797	65.303	1.00 14.70	SOLV
HETATM	3021	OH2	WAT	ם	12	20.490	34.049	61.067	1.00 25.01	SOLV
HETATM			WAT		13	44.757	33.106	46.084	1.00 25.90	SOLV
HETATM	3022								1.00 15.21	SOLV
HETATM			WAT		14	22.457	60.823	57.444		
HETATM	3024		WAT		15	3.399	32.742	65.163	1.00 20.66	SOLV
HETATM	3025	OH2	WAT	D	16	32.273	51.414	45.610	1.00 22.37	SOLV
HETATM		OH2	WAT	D	17	26.328	42.873	73.427	1.00 27.86	SOLV
HETATM	3027		WAT		18	48.249	24.121	56.778	1.00 15.09	SOLV
HETATM	3038		WAT		19	15.249	44.552	72.082	1.00 40.95	SOLV
MENTAN	3040						9.269	52.633	1.00 26.66	SOLV
HETATM	3049	OH2	WAT		20	26.444				
HETATM	3030	OH2	WAT		21	26.554	18.383	59.650	1.00 11.42	SOLV
HETATM	3031	OH2	TAW	D	22	39.456	25.964	72.316	1.00 20.32	SOLV
HETATM	3032	OH2	WAT	D	23	26.743	37.600	38.359	1.00 37.22	SOLV
HETATM	3033	OH2	WAT		24	44.666	23.818	39.068	1.00 32.27	SOLV
HETATM	3034		WAT			14.714	52.213	70.663	1.00 29.24	SOLV
HETATM	3034				26	45.129		69.864	1.00 29.58	SOLV
HETATM	3035	OHZ	TAW	ב			18.856			
HETATM	3036	OH2	WAT	מ	27	30.024	17.886	49.758	1.00 15.52	SOLV

HETATM	3037	OH2	WAT	D	28	20.659	28.788	43.520	1.00 28.55	SOLV
HETATM	3038	OH2	WAT	D	29	32.271	38.000	53.512	1.00 47.72	SOLV
HETATM	3039	OH2	WAT	D	30	18.285	29.333	54.536	1.00 21.34	SOLV
HETATM	3040	OH2	WAT	D	31	49.978	38.669	73.461	1.00 31.02	SOLV
HETATM	3041	OH2	WAT	D	32	21.587	50.386	71.043	1.00 14.52	SOLV
HETATM	3042	OH2	WAT	D	33	46.784	32.121	33.375	1.00 31.79	SOLV
HETATM			WAT		34	33.359	39.755	49.117	1.00 16.13	SOLV
HETATM			WAT		35	7.687	37.657	51.568	1.00 27.22	SOLV
HETATM			WAT		36	44.238	35.392	33.961	1.00 19.67	SOLV
HETATM			WAT		37	10.908	25.384	58.206	1.00 33.51	SOLV
HETATM			WAT		38	36.758	27.243	70.552	1.00 39.61	SOLV
HETATM			WAT		39	45.825	46.691	54.654	1.00 32.43	SOLV
. HETATM			WAT		40	52.489	20.282	52.165	1.00 39.37	SOLV
HETATM			WAT		42	12.117	17.831	56.596	1.00 27.74	SOLV
HETATM			WAT		. 43	45.023	26.168	35.172	1.00 27.74	
			WAT		44	39.392	12.771	62.066		SOLV
HETATM			WAT			3,930	26.970		1.00 35.15	SOLV
HETATM					45			63.814		SOLV
HETATM			WAT		46	8.454	19.321	71.677	1.00 32.36	sorv
HETATM			WAT		47	20.280	18.126	73.237	1.00 33.88	SOLV
HETATM			WAT		48	9.321	39.409	54.873	1.00 18.57	SOLV
HETATM			WAT		49	50.852	41.323	58.048	1.00 21.25	SOLV
HETATM			WAT		50	37.134	34.599	60.315	1.00 61.70	SOLV
HETATM			WAT		- 51	14.944	62.815	48.613	1.00 42.50	SOLV
HETATM			WAT		52	6.494	33.164	51.420	1.00 40.65	SOLV
HETATM			TAW		53	24.913	44.799	72.298	1.00 17.10	SOLV
HETATM			WAT		54	51.156	35.095	48.814	1.00 23.05	SOLV
HETATM			WAT		55	16.518	41.750	45.596	1.00 49.25	SOLV
HETATM			WAT		56,	10.326	16.413	61.267	1.00 46.03	SOLV
HETATM			WAT		57	25.316	47:708	73.062	1.00 22.73	SOLV
HETATM			WAT		58	4.013	33.865	76.173	1.00 44.82	SOLV
HETATM			WAT		59	24.846	18.072	36.805	1.00 34.67	SOLV
HETATM			WAT		60	15.930	56.853	61.737	1.00 55.56	SOLV
HETATM			WAT		61	49.662	44.249	48.982	1.00 28.72	SOLV
HETATM			WAT		62.	23.232	17.421	53.920	1.00 13.11	SOLV
HETATM			WAT		63	39.293	23.035	33.289	1.00 35.79	SOLV
HETATM			WAT		64	19.908	20.169	44.339	1.00 24.33	SOLV
HETATM			TAW		65	33.259	21.655	69.560	1.00 45.10	SOLV
HETATM			TAW		66	27.528	53.947	68.629	1.00 44.79	SOLV
HETATM			TAW		67	18.774	48.716	52.865	1.00 54.01	SOLV
HETATM	_		TAW		68	10.877	29.062	63.401	1.00 27.08	SOLV
HETATM			TAW		69	43.057	31.367	28.786	1.00 30.16	SOLV
HETATM			WAT		70	24.816	44.057	43.447	1.00 20.11	SOLV
HETAŢM			WAT		71	37.368	38.823	46.381	1.00 33.55	SOLV
HETATM			TAW		72	9.038	18.327	63.519	1.00 31.34	SOLV
HETATM			TAW		73	51.799	20.829	65.265	1.00 28.32	· SOLV
HETATM			WAT		74	17.556	58.515	57.254	1.00 19.27	SOLV
HETATM			WAT		75	28.436	27.904	79.425	1.00 27.13	SOLV
HETATM			WAT		76	18.939	35.798	35.800	1.00 94.18	SOLV
HETATM			WAT		77	34.359	31.251	46.688	1.00 73.70	SOLV
HETATM			WAT		78	44.373	51.649	60.029	1.00 30.23	SOLV
METATM	3087		WAT		79	28.537	63.478	48.324	1.00 21.09	SOLV
HETATM	3088	OH2	WAT	D	80	6.869	44.113	72.030	1.00 28.59	SOLV
HETATM	3089		WAT		81	42.882	18.761	71.115	1.00 31.80	SOLV
HETATM	3090	OH2	TAW	D	82	36.712	59.078	53.901	1.00 40.11	SOLV
HETATM	3091	OH2	TAW	D	83	37.506	42,495	40.104	1.00 51.37	SOLV
HETATM	3092	OH2	TAW	D	84	40.054	38.439	55.415	1.00 20.07	SOLV
HETATM	3093	OH2	WAT	D	85	32.170	56.633	72.920	1.00 45.23	SOLV
HETATM		OH2	WAT	D	86	24.470	53.877	47.119	1.00 41.18	SOLV
HETATM	3095		WAT		87	48.585	35.663	67.518	1.00 33.40	SOLV
HETATM			WAT		88	29.541	57.166	42.788	1.00 44.61	SOLV
HETATM			WAT		89	47.814	28.707	41.228	1.00 45.64	SOLV
HETATM			WAT		90	49.377	52.112	63.320	1.00 22.26	SOLV
HETATM	3099		WAT		91	44.219	43.589	43.912	1.00 39.90	SOLV
HETATM	3100		WAT		92	25.913	61.639	75.382	1.00 48.28	SOLV
HETATM	3101	OHO	WAT	ח	93	8.623	30.749	49.707	1.00 40.37	SOLV
HETATM			WAT		94	45.634	41.080	40.990	1.00 21.46	SOLV
ELIAIM	2102	V.12		_						

### SUBSTITUTE SHEET (RULE 26)

HETATM	3103	OH2	WAT	D	95		29.984	34.886	51.725	1.00 3	5.75	SOLV
HETATM		OH2	TAW	D	96		13.051	21.934	49.804	1.00 4	6.73	SOLV
HETATM		OH2	WAT	D	97		32.412	65.913	55.822	1.00 4	3.39	SOLV
HETATM		OH2	TAW	D	98		35.056	43.390	38.348	1.00 3	4.53	SOLV
HETATM			TAW		99		22.360	47.680	60.688	1.00 1	9.16	SOLV
HETATM			WAT				50.755	19.722	57.906	1.00 4	2.45	SOLV
HETATM	3100		WAT				7.875	37.690	74.094	1.00 3		SOLV
HETATM	3110				102		24.080	26.796	43.617	1.00 3		SOLV
			WAT				45,206	34.126	75.765	1.00 3		SOLV
HETATM			WAT				26.110	54.786	40.685	1.00 2		SOLV
HETATM	3112		WAT				25.918	39.658	77.647	1.00 4		SOLV
HETATM						-	41.578	18.191	36.809	1.00 4		SOLV
HETATM			TAW				31.945	51.420	73.896	1.00 4		SOLV
HETATM			WAT									
HETATM			WAT	•			16.722	60.311	51.182	1.00 4		SOLV
HETATM			WAT				43.604	38.573	78.141	1.00 3		SOLV
HETATM			WAT				16.063 21.630	15.496	69.430	1.00 5		SOLV
HETATM	3119		WAT				21.630	22.785	49.145	1.00 3		SOLV
HETATM	3120		TAW				27.479	56.647	44.026	1.00 5		soŕv
HETATM	3121	OH2	TAW	D	113		14.739	51.674	61.674	1.00 3		SOLV
HETATM	3122	OH2	WAT	D	114		50.063	26.435	54.358	1.00 5		SOLV
HETATM	3123	OH2	WAT	D	115		43.935	38.427	73.129	1.00 4		SOLV
HETATM	3124	OH2	TAW	D	116		49.707	31.478	57.709	1.00 3		SOLV
HETATM	3125	OH2	WAT	D	117		25.032	43.463	55.676	1.00 3		
HETATM		OH2	WAT	D	118		10.618	46.623	59.838	1.00 2		SOLV
HETATM	3127	OH2	WAT	D	119		48.466	33.382	61.437	1.00 1	9.82	SOLV
HETATM		OH2	WAT	D	120		44.157	40.058	.37.907	1.00 4	2.95	SOLV
HETATM		OH2	WAT	D	121		51.267	29.446	52.889	1.00 3		SOLV
HETATM		OH2	WAT	D	122		16.653	15.228	72 ⁻ .975	1.00 4	5.41	SOLV
HETATM		OH2	TAW	D	123	•	36.898	45.148	41.936	1.00 2	7.00	SOLV
HETATM		OH2	WAT	D	124		49.655	34.591	5 <b>9.117</b>	1.00 3	8.97	SOLV
HETATM		OH2	TAW	D	125		12.285	57.594	42.107	1.00 2	3.56	SOLV
HETATM		OH2	WAT	D	126		28.294	57.644	73.289	1.00 3	4.79	SOLV
HETATM			WAT				19.138	60.403	61.551	1.00 2	8.58	SOLV
HETATM			WAT				30.300	33.685	34.047	1.00 2	7.37	SOLV
HETATM			WAT				40.898	53.983	47.254	1.00 1	6.30	SOLV
HETATM			WAT				43.550	32.160	38.272	1.00 3	8.86	SOLV
HETATM			WAT				18.624	13.959	56.194	1.00 3	7.70	SOLV
HETATM			WAT				18.580	12.901	62.894	1.00 2	7.28	SOLV
HETATM			WAT				35.830	30.296	50.621	1.00 4	2.47	SOLV
HETATM		OH2	WAT	D	134		51.219	35.855	51.878	1.00 2	0.37	SOLV
HETATM			WAT				50.428	22.486	49.267	1.00 3	9.37	SOLV
HETATM			WAT				51.633	29.369	63.918	1.00 3		SOLV
HETATM			WAT				46.384	43.924	55.825	1.00 2		SOLV
			WAT				30.356	25.767	28.762	1.00 2		SOLV
HETATM HETATM	3147		TAW				25.070	47.842	60.819	1.00 2		SOLV
HETAIM	31/0		WAT				47.097	49.394	69.367	1.00 3		SOLV
HETATM	3140	202	WAT	ח	141		15.246	37.581	73.398	1.00 3		SOLV
HETATM	3149	727	WAT	ב	142		8.341	23.099	64.695	1.00 3		SOLV
HETATM		2.112	WAT	5	142		30.065	18.220	46.048	1.00 1		SOLV
HETATM		Zn'y	WAT	7	144		11.930	46.453	57.606	1.00 3		SOLV
HETATM	3152	OHZ	MW.T.	יי	764		11.320	40.403	57.000	4.00 3	0.13	2010

#### INTERNATIONAL SEARCH REPORT

International application No. PCT/US00/24700

A. CLASSIFICATION OF SUBJECT MATTER  IPC(7) :CO7K 14/00; GO1N 53/573  US CL :Please See Extra Sheet.  According to International Patent Classification (IPC) or to both national classification and IPC  B. FIELDS SEARCHED  Minimum documentation searched (classification system followed by classification symbols)  U.S. : Please See Extra Sheet.								
Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched								
Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)  Please See Extra Sheet.								
C. DOCUMENTS CONSIDERED TO BE RELEVANT								
Category*	Citation of document, with indication, where ap	propriate, of the relevant passages	Relevant to claim No.					
J	KAKUTA et al. Crystal Structure of the Human Heparan Sulfate N-Deacetylas Journal of Biological Chemistry. 16 Number 16, pages 10673-10676, see et	se/N-Sulfotransferase 1. The April 1999, Volume 274,	1-19					
o I	SUEYOSHI et al. A role of Lys-614 in a function of human heparan sulfate N-deacetula Letters. 1998, Volume 433, pages in a stract.	se/N-sulfotransferase. FEBS	1-19					
X Further documents are listed in the continuation of Box C. See patent family annex.								
	l extegories of cited documents: ent defining the general state of the art which is not considered	"I" later document published after the inter date and not in conflict with the appli	cation but cited to understand					
to be a	of particular relevance  character the international filing date	the principle or theory underlying the invention  "X" document of particular relevance; the claimed invention cannot be						
"L" docume	ent which may throw doubts on priority claim(s) or which is to establish the publication date of another citation or other	considered novel or cannot be considered to involve an inventive step when the document is taken alone						
special	ent referring to an oral disclosure, use, exhibition or other	"Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being						
means		obvious to a person skilled in the art  document member of the same patent family						
than ()	he priority date claimed tual completion of the international search	Date of mailing of the international search report						
	29 DECEMBER 2000 25 JAN 2001							
	ling address of the ISA/US of Patents and Trademarks  O.C. 20231 (703) 305-3230	Authorized officer  ARDIN MARSCHEL  Telephone No. (708) 508-0196						

#### INTERNATIONAL SEARCH REPORT

International application No. PCT/US00/24700

	16176.	3007 24700
C (Continua	ntion). DOCUMENTS CONSIDERED TO BE RELEVANT	
Category*	Citation of document, with indication, where appropriate, of the relevant passag	es Relevant to claim No
Y	AHMAD et al. WD Repeats of the p48 Subunit of Chicken Chromatin Assembly Factor-1 Required for in Vitro Interaction with Chicken Histone Deacetylase-2. The Journal of Biologica Chemistry. 04 June 1999, Volume 274, Number 23, pages 16646 16653, see especially the abstract.	1
Y	JOHN et al. Rhizobium NodB protein involved in nodulation signal synthesis is a chitooligosaccharide deacetylase. Proceedi of the National Academy of Sciences, USA. January 1993, Vol 90, pages 625-629, see especially the abstract.	ngs
A	US 5,780,594 A (CARTER) 14 July 1998, see the entire disclose	ure. 1-19
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#### INTERNATIONAL SEARCH REPORT

International application No. PCT/US00/24700

A. CLASSIFICATION OF SUBJECT MATTER: US CL  $\,:\,$ 

530/350 and 435/7.9

B. FIELDS SEARCHED
Minimum documentation searched
Classification System: U.S.

530/300,333,350; 435/6,7.2; 514/2

**B. FIELDS SEARCHED** 

Electronic data bases consulted (Name of data base and where practicable terms used):

CAS, BIOTECH ABS, MEDLINE, EMBASE, WPI, WEST covering search terms: deacetylase, human, crystal, histone, inhibitor, x-ray, and crystallography

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